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2 09:05:31 2003
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GenCore version 5.1.6
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OM protein - protein search, using sw model

December 2, 2003, 02:15:49 ; Search time 218.096 Seconds Run on:

(without alignments) 246.943 Million cell updates/sec

US-09-607-745-9

1600 1 MDSKGSSQKSRLLLLLVVSN......LNWIYNVWKAELSRHHHHHH 292 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

Total number of hits satisfying chosen parameters: 673684 seqs, 184443283 residues

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Published Applications AA:*

1: /cgn2_6/ptodata/2/pubpaa/US07 PUBCOMB.pep:*

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/cgn2_6/ptodata/2/pubpaa/US60_PUBGOMB.pep:* ptodata/2/pubpaa/US10B_PUBCOMB.pep ptodata/2/pubpaa/US10C

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		Appl 1	72. April	23, Appl		Appli	275. Ann				330, Ann		330. App			330, App	
	Description	Sequence 6. Appli	Segmence 7	Sequence 2		Semience 7											
	ID	US-09-851-588-6	US-09-776-191-72	US-10-264-820-23	US-10-254-289~2	US-09-888-257A-7	US-09-946-374-275	US-10-015-387A-275	US-10-063-735-112	US-10-006-130A-275	US-10-199-672-330	US-10-006-172A-275	US-10-187-749-330	US-10-194-457-330	US-10-184-642-330	US-10-196-747-330	
	DB	6	11	15	15	11	11	12	12	12	12	12	12	12	12	12	
	(atch Length DB	406	423	423	423	432	432	432	432	432	432	432	432	432	432	432	
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	Score	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	1266	
Result	No.	1	7	m	4	D.	9	7	80	თ	10	11	12	13	14	15	

111 KLGSFPSLAVAKIIIIBFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEELTPATP 170

233 KLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEELTPATP

173 RVVGGERASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSD

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171 LWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAYQGEVTEKMMCAGIPEGGVDTC 230

App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App	App
275,	275,	33	33	33	33	3	33	33	m	33	33	33	33	33	33	33	33	33	33	33	3	33	33	33	33	330,	330,	330,	330,
Sequence	Sequence	Sequence	Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Seguence	Sequence	Sequence	Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Seguence	Seguence
-015-392A-2	-10-017-253A-27	-173 - 689 - 33	-10-173-690-33	-10-173-691-33	-173-692-	-10-173-694-33	-10-173-698-33	0-173-699-33	-10-173-	-10-174-569-33	-10-174-583-33	-174-587-33	-10-174-589-33	-10-174-591-33	-10-175-736-33	-10-175-742-33	-10-175-744-33	-10-175-745-33	-10-175-748-33	10-175-751-33	-10-175-754-33	10-176-480-33	-10-176-489-33	10-176-754-33	-10-176-755-33	-10-176-759-33	-10-176-920-	-10-176-922-33	4-33
12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	77	12	12		12	12		12	12	12	12	17	7,5
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16	17	18	19	20	21	22	23	24	25	56	27	28	29	30	31	32	33	34	32	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Sequence 6, Application US/09851588

Patent No. US2002042067A1

GENERAL INFORMATION:

APPLICANT: Mack, David

APPLICANT: Gish, Kurt C.

TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AN.

TITLE OF INVENTION NUMBER: US 09/656,002

PRIOR APPLICATION NUMBER: US 09/656,002

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PARCENT IN VERSION 3.1

FRANCE FILING DATE: 2000-09-06

NUMBER OF SEQ ID NOS: 9

SOFTWARE: PARCENT IN VERSION 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51 KIVGGYALDVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 79.1
Best Local Similarity 97.4
Matches 228; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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US-09-851-588-6
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190 RVVGGEBASVDSWPWQVSIQYDKQHVCGGSILDPHWVJTAAHCFRKHTDVFNWKVRAGSD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     310 LWIIGWGFTKONGGKMSDILLQASVQVIDSTRCNADDAYQGEVTEKMMCAGIPEGGVDTC
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Publication No. US20030118509A1
GGRERAL INFORMATION:
APPLICANT: Wack, David
APPLICANT: Wilson, Keith
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMF
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMF
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMF
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMF
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMF
FILE REFERENCE: A-69108/DJD/AMS
CURRENT APPLICATION NUMBER: US/09/656,002
PRIOR APPLICATION NUMBER: US/09/656,002
PRIOR PILING DATE: 2000-09-06
PRIOR PRILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-01-28
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
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2; Mismatches
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 09/268,866
PRIOR PILING DATE: 1999-03-15
PRIOR APPLICATION NUMBER: US 09/435,945
PRIOR PILING DATE: 1999-11-09
PRIOR PILING DATE: 1999-11-09
PRIOR PILING DATE: 1999-11-09
PRIOR PILING DATE: 1999-11-29
PRIOR PILING DATE: 1999-11-29
PRIOR PILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: US 09/453,850
PRIOR PILING DATE: 1999-12-02
PRIOR PILING DATE: 1999-12-02
PRIOR PILING DATE: 1999-12-03
PRIOR PILING DATE: 1999-12-03
PRIOR PILING DATE: 2000-01-28
PRIOR PILING DATE: 2000-03-15
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Best Local Similarity 97.4%;
Matches 228; Conservative
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SOFWHARE: Patentin version 3.0
SEQ ID NO .
ERNOTH: 423
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 23
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Publication No. US20030108926A1
GENERAL INFORMATION: No. US20030108926A1e1 Methods of Diagnosing Colorectal Cancer,
TITLE OF INVENTION: Compositions, and Methods of Screening for Colorectal
TITLE OF INVENTION: Cancer Modulators
FILE REFERENCE: 018501-006141US
CURRENT APPLICATION NUMBER: US/10/264,820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Edgar O. Ong
APPLICANT: Edgar O. Ong
APPLICANT: Edgar O. Ong
APPLICANT: Junn-Chern Yeh
APPLICANT: Junn-Chern Yeh
APPLICANT: Ocras International, Inc.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING
TITLE OF INVENTION: METHODS BASED THEREON
TILLE OF INVENTION NUMBER: 60/213,124
PRIOR APPLICATION NUMBER: 60/213,124
PRIOR APPLICATION NUMBER: 60/213,4840
PRIOR PILING DATE: 2000-06-22
PRIOR PILING DATE: 2000-06-22
PRIOR FILING DATE: 2000-02-08
PRIOR FILING DATE: 2000-02-03
PRIOR FILI
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        231 QGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKABL 284
                                                                                                                                                                             353 QGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKAEL 406
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                                                                                                                                                                                                                                                                                                                      RESULT 2
US-09-776-191-72
US-09-776-191-72
; Publication No. US20030119168A1
; GENERAL INFORMATION:
; APPLICANT: Edwin L. Madison
; APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: A
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CRGANISM: Homo sapien
US-09-776-191-72
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COMPOSITIONS,

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259 KLGSFPSLAVAKIIIIEFNPWYPKDNDIALMKLQPPLTFSGTVRPICLPFFDEELTPATP
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                                                                                                                                    Length 432;
                                                                                                                                                                                                  Indels
                                                                                                                                    Score 1266; DB 11;
Pred. No. 6.3e-117;
                                                                                                                                                                                               2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, Micholas F.
APPLICANT: Stewart, Timothy A.
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Watliams, P. Mickey
APPLICANT: Watliams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrar
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2380PLG1
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2380PLG1
CURRENT APPLICATION NUMBER: G0/098716
PRIOR APPLICATION NUMBER: 60/098716
PRIOR APPLICATION NUMBER: 60/098729
PRIOR APPLICATION NUMBER: 60/098749
PRIOR APPLICATION NUMBER: 60/098749
PRIOR APPLICATION NUMBER: 60/098749
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/09834
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/09834
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/09834
PRIOR FILING DATE: 1998-09-02
PRIOR APPLICATION NUMBER: 60/09936
PRIOR APPLICATION NUMBER: 60/09936
PRIOR APPLICATION NUMBER: 60/09936
PRIOR APPLICATION NUMBER: 60/09936
PRIOR PRILING DATE: 1998-09-09
PRIOR APPLICATION NUMBER: 60/09936
PRIOR APPLICATION NUMBER: 60/09936
PRIOR PRILING DATE: 1998-09-09
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Publication No. US20030073129A1
GENERAL INFORMATION:
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Grimaldi, Christopher J.
Gurney, Austin L.
Hillan, Kenneth J.
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Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
                                                                                                                              Query Match
Best Local Similarity 97.4
Matches 228; Conservative
                                      ; ORGANISM: Homo Sapien
US-09-888-257A-7
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US-09-946-374-275
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APPLICANT:
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                                                                                                                                                                                                                                         190 RVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSD
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
FILE REPERENCE: PSOOZH
CURRENT APPLICATION NUMBER: US (9/08),635
PRIOR APPLICATION NUMBER: US (6/08),540
PRIOR PILING DATE: 1997-10-28
PRIOR PILING DATE: 1997-10-28
PRIOR FILING DATE: 1998-06-17
PRIOR PILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-17
PRIOR PILING DATE: 1998-06-10-08
PRIOR FILING DATE: 1999-06-01
PRIOR PILING DATE: 2000-03-18
PRIOR PILING DATE: 2000-03-18
PRIOR PILING DATE: 1999-06-01
PRIOR PILING DATE: 2000-03-18
PRIOR PILING DATE: 2001-03-01
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                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          370 QGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKAEL 423
                                                                                                                                 0
                                                             Query Match
79.1%; Score 1266; DB 15; Length 423;
Best Local Similarity 97.4%; Pred. No. 6.2e-117;
Matches 228; Conservative 2; Mismatches 4; Indels 0
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Publication No. US20030060612A1
GENERAL INFORMATION:
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
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Smith, Victoria
US-10-254-289-2
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PRIOR FILING DATE: 1998-09-09
PRIOR PELICATION UNDER: 60/09508
PRIOR APPLICATION UNDER: 60/099642
PRIOR PELICATION UNDER: 60/099642
PRIOR PELICATION UNDER: 60/099642
PRIOR FILING DATE: 1998-09-09
PRIOR PELICATION UNDER: 60/099741
PRIOR FILING DATE: 1998-09-10
PRIOR PELICATION UNDER: 60/09972
PRIOR FILING DATE: 1998-09-10
PRIOR PELICATION UNDER: 60/09972
PRIOR PELICATION UNDER: 60/099012
PRIOR PELICATION UNDER: 60/109912
PRIOR PELICATION UNDER: 60/100961
PRIOR PELICATION UNDER: 60/100961
PRIOR PELICATION UNDER: 60/100661
PRIOR PELICATION UNDER: 60/10061
PRIOR PELICATION UNDER: 60/10010
PRIOR PELICATION UNDER: 6

PRIOR APPLICATION NUMBER: 60/101479
PRIOR FILING DATE: 1998-09-23
PRIOR FILING DATE: 1998-09-23
PRIOR FILING DATE: 1998-09-24
PRIOR PRILING DATE: 1998-09-29
PRIOR APPLICATION NUMBER: 60/101916
PRIOR APPLICATION NUMBER: 60/101916
PRIOR APPLICATION NUMBER: 60/10240
PRIOR PRILING DATE: 1998-09-29
PRIOR PILING DATE: 1998-09-30
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/103395
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/103495
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/103495
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/103495
PRIOR PILING DATE: 1998-10-08
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/103499
PRIOR PILING DATE: 1998-10-08
PRIOR PILING DATE: 1998-10-08
PRIOR PILING DATE: 1998-10-07
PRIOR PILIN

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Ferrara, Napoleone
Fong, Sherman
Gao, Wei-Qiang
                  171 LWIIGWGFTKONGGKMSDI
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Goddard, Audrey
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan 1.
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Matches 228; Conservative
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ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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US-10-063-735-112
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            319 LWIIGWGFTKONGGKMSDILLQASVQVIDSTRCNADDAYQGEVTBKWMCAGIPEGGVDTC 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids ancoding the Same FILE REFERENCE: P2830PLC54
CURRENT APPLICATION NUMBER: US/10/015,387A
CURRENT FILING DATE: 2001-12-12
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 477
SEQ ID NO 275
LENGTH: 432
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                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                           79.1%; Score 1266; DB 11; Length 432;
larity 97.4%; Pred. No. 6.3e-117;
Conservative 2; Mismatches 4; Indels 0
                                                                                                                                                                                              Pred. No. 6.3e-117
2; Mismatches 4
              PRIOR APPLICATION NUMBER: 60/105266
PRIOR FILING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: 60/105693
PRIOR FILING DATE: 1998-10-26
PRIOR APPLICATION NUMBER: 60/105694
PRIOR PILING DATE: 1998-10-26
PRIOR APPLICATION NUMBER: 60/105807
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Fong, Sherman
Gao, Wei-Qian
Godoard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
Gurney, Austin L.
Hillan, Kenneth J.
Pan, James
Paoni, Nicholas F.
PRIOR FILING DATE: 1998-10-22
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan 1.
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Best Local Similarity 97.4
Matches 228; Conservative
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, ORGANISM: Homo sapiens
US-10-015-387A-275
                                                                                                                                                                                              Best Local Similarity
Matches 228; Conserv
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APPLICANT:
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230
                                                          319 LWIIGWGFTKONGGKASDILLQASVOVIDSTRCNADDAYQGEVTEKMMCAGIFEGGVDTC 378
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurnery, Austrian L.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMENANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P2330R1C1
CURRENT APPLICATION NUMBER: US/10/063,735
CURRENT FILING DATE: 2002-05-08
Prior Application removed - See Palm or File Wrapper NUMBER OF SEQ ID NOS: 170
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                                                                                                                                                                                         379 QGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKAEL 432
                                                                                                                                  QGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKAEL
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                                                                                                                                                                                                                                                                                                                                                       Sequence 112, Application US/10063735
Publication No. US20030138882A1
GENERAL INFORMATION:
APPLICANT: Eaton, Dan L.
APPLICANT: Filvaroff, Ellen
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Godowski, Paul J.
Grimaldi, Christopher J.
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APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan 1.
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Best Local Similarity 97.4
Matches 228; Conservative
                                                                                                                                                                                                                                                       ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-199-672-330
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APPLICANIT: AMONO, MILLIAM I.

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC

TITLE OF INVENTION: ACIDS ENCODING THE SAME

TITLE OF INVENTION: ACIDS ENCODING THE SAME

TITLE OF INVENTION: ACIDS ENCODING THE SAME

TITLE OF INVENTION NUMBER: US/10/199,672

CURRENT APPLICATION NUMBER: US/10/052,586

PRIOR APPLICATION NUMBER: 60/059263

PRIOR APPLICATION NUMBER: 60/059266

PRIOR APPLICATION NUMBER: 60/059266

PRIOR APPLICATION NUMBER: 60/062250

PRIOR APPLICATION NUMBER: 60/062120

PRIOR APPLICATION NUMBER: 60/063121

PRIOR APPLICATION NUMBER: 60/063121

PRIOR FILING DATE: 1997-10-24

PRIOR APPLICATION NUMBER: 60/063121

PRIOR APPLICATION NUMBER: 60/063121

PRIOR FILING DATE: 1997-10-24

PRIOR FILING DATE: 1997-10-24

PRIOR FILING DATE: 1997-10-24

PRIOR FILING DATE: 1997-10-24
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APPLICANT: Pan, James
APPLICANT: Pan, James
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2830PIC7
CURRENT APPLICATION NUMBER: US/10/006,130A
CURRENT FILING DATE: 2002-03-19
NUMBER OF SEQ ID NOS: 477
SEQ ID NO 275
LEMANT. . . .
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Pred. No. 6.3e-117;
2; Mismatches 4;
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p. US20030148442A1
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Watanabe, Colin K.
Wood, William I.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 97.45
Matches 228; Conservative
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                                                                                                                                                                                                                                                                                                                   TYPE: PRT
CORGANISM: Homo sapiens
US-10-006-130A-275
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Publication No. US20
GENERAL INFORMATION:
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APPLICANT:
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51 KIVGGYALDVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSD 110
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APPLICANT: Hillan, Kenneth J.
APPLICANT: Pan James
APPLICANT: Pani, Usicholas F.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
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PRIOR APPLICATION NUMBER: 60/063540
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063541
PRIOR FILING DATE: 1997-10-28
PRIOR FILING DATE: 1997-10-28
PRIOR PILING DATE: 1997-10-28
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 330
LENGTH: 432
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Pred. No. 6.3e-117;
2; Mismatches 4;
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CURRENT APPLICATION NUMBER: US/10/006,172A
CURRENT APPLICATION NUMBER: US/10/006,172A
CURRENT FILING DATE: 2002-03-19
PRIOR APPLICATION NUMBER: 60/098716
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR FILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR APPLICATION NUMBER: 60/098760
PRIOR PILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098803
PRIOR FILING DATE: 1998-09-02
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Fong, Sherman
Gao, Wei-Ciang
Goddard, Audrey
Goddwski, Paul J.
Grimaldi, Christopher J.
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NUMBER: 6 1998-09- NUMBER: 6 1998-09- NUMBER: 6 1998-09- NUMBER: 6 1998-09- NUMBER: 6 1998-09- NUMBER: 6 1998-09-	NUMBER: 60 1998-09-1 1998-09-1 1998-09-1 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60 1998-09-1 NUMBER: 60	1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1 1998-09-1	1998-09-18 NUMBER: 60/10084 1998-09-18 NUMBER: 60/10091 1998-09-17 NUMBER: 60/10091 1998-09-17 NUMBER: 60/10101 1998-09-18 NUMBER: 60/10107 1998-09-18 NUMBER: 60/10107 1998-09-18 NUMBER: 60/10127 1998-09-18 NUMBER: 60/10147 1998-09-23 NUMBER: 60/10147 1998-09-23 NUMBER: 60/10147 1998-09-23 NUMBER: 60/10147
APPLICATION APPLICATION APPLICATION FILING DATE: APPLICATION APPLICATION FILING DATE: APPLICATION FILING DATE: APPLICATION	APPLICATION APPLICATION APPLICATION FILING DATE: FILING DATE: APPLICATION APPLICATION FILING DATE: APPLICATION FILING DATE APPLICATION	PELLING DATE PILING DATE PILING DATE APPLICATION PILING DATE APPLICATION PILING DATE APPLICATION PILING DATE APPLICATION PILING DATE PILING DATE PILING DATE PILING DATE	APPLICATIONS APPLI
PRIOR	PRIOR PRIOR	PRIOR	PRICE PRICE

PRIOR FILING DATE: 1998-09-23
PRIOR PELICATION WINDERS: 60/101479
PRIOR APPLICATION WINDERS: 60/101473
PRIOR APPLICATION WINDERS: 60/101473
PRIOR APPLICATION WINDERS: 60/101413
PRIOR APPLICATION WINDERS: 60/101413
PRIOR APPLICATION WINDERS: 60/101413
PRIOR PLING DATE: 1998-09-24
PRIOR PLING DATE: 1998-09-29
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-30
PRIOR PLING DATE: 1998-10-07
PRIOR PLING DATE: 1998-10-

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TYPE: PRT; ORGANISM: Homo Sapien
US-10-187-749-330
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CURRENT APPLICATION NUMBER: US/10/052,586

PRIOR APPLICATION NUMBER: US/10/052,586

PRIOR FILING DATE: 2002-01-15

PRIOR FILING DATE: 1997-09-18

PRIOR APPLICATION NUMBER: 60/059266

PRIOR FILING DATE: 1997-09-18

PRIOR FILING DATE: 1997-01-17

PRIOR FILING DATE: 1997-01-17

PRIOR FILING DATE: 1997-01-17

PRIOR FILING DATE: 1997-10-17

PRIOR FILING DATE: 1997-10-17

PRIOR FILING DATE: 1997-10-17

PRIOR FILING DATE: 1997-10-24

PRIOR FILING DATE: 1997-10-24
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PRIOR FILING DATE: 1998-10-21
PRIOR APPLICATION NUMBER: 60/105169
PRIOR PLING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: 60/105169
PRIOR PLING DATE: 1998-10-22
PRIOR PILING DATE: 1998-10-22
PRIOR PELING DATE: 1998-10-26
PRIOR FILING DATE: 1998-10-26
PRIOR FILING DATE: 1998-10-26
PRIOR PLING DATE: 1998-10-26
PRIOR APPLICATION NUMBER: 60/105807
PRIOR APPLICATION NUMBER: 60/105807
PRIOR PRIING DATE: 1998-10-27
PRIOR PULING DATE: 1998-10-27
PRIOR PLING DATE: 1998-10-27
PRIOR PLING DATE: 1998-10-27
PRIOR PILING DATE: 1998-10-27
PRIOR APPLICATION NUMBER: 60/105801
PRIOR PILING DATE: 1998-10-27
PRIOR APPLICATION NUMBER: 60/106023
PRIOR PILING DATE: 1998-10-27
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o. US20030153036A1
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Pan, James
Smith, Victoria
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Publication No. US20
GENERAL INFORMATION:
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APPLICANT:
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APPLICANT:
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APPLICANY: Goddard, Audrey
APPLICANY: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Gurney, Austin L.
APPLICANT: Bu, James
APPLICANT: Smith, Victoria
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: BECRETED AND TRANSNEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION NUMBER: 00/059263
PRIOR FILING DATE: 1997-09-18
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/063120
PRIOR FILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063121
PRIOR FILING DATE: 1997-10-24
PRIOR FILING DATE: 1997-10-21
PRIOR FILING DATE: 1997-10-21
PRIOR FILING DATE: 1997-10-21
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PRIOR APPLICATION NUMBER: 60/063540
PRIOR FILING DATE: 1997-10-21
PRIOR APPLICATION NUMBER: 60/063540
PRIOR PLING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063541
PRIOR FILING DATE: 1997-10-28
REMEMBRING PRIOR APPLICATION NUMBER: 60/063544
PRIOR FILING DATE: 1997-10-28
REMEMBRING PRIOR APPLICATION DATE: 1997-10
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Publication No. US20030153037A1
GENERAL INFORMATION:
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US-10-196-747-330
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TITLE OF INVENTION: SECRETED AND TRANSMENBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3430RIC194
CURRENT APPLICATION NUMBER: US/10/184,642
CURRENT FILING DATE: 2002-06-27
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Pred. No. 6.3e-117;
2; Mismatches 4; Indels 0
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                    PRIOR APPLICATION NUMBER: 60/063541
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: 60/063544
PRIOR PILING DATE: 1997-10-28
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SEQ ID NO 330
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Publication No. US20030157635A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
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97.4%;
PRIOR FILING DATE: 1997-10-28
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Godowski, Paul J.
Gurney, Austin L.
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Best Local Similarity 97.4 Matches 228, Conservative
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Smith, Victoria
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ORGANISM: Homo Sapien
US-10-184-642-330
                                                                                                                                                                                                   TYPE: PRT; ORGANISM: Homo Sapien
US-10-194-457-330
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259 KLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEELTPATP 318
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PLIAND APPLICATION NUMBER: US/10/196,747
Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 612
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Pred. No. 6.3e-117;
2; Mismatches 4;
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Publication No. US20030162250Al
GENERAL INFORMATION:
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Best Local Similarity 97.4%;
Matches 228; Conservative
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Wood, William I.
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Gurney, Austin L.
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Goddard, Audrey
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Copyright (c) 1993 - 2003 Compugen Ltd.
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Listing first 45 summaries
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RESULT 2 US-09-008-271A-6

Appli Appli Appli

Sequence 3, Appli	Sequence 26, Appl	Sequence 63, Appl	Sequence 44, Appl	Sequence 68, Appl	Sequence 4, Appli		Sequence 7, Appli	Sequence 2, Appli	'n	Sequence 4, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 13, Appl	Sequence 7, Appli	H	Sequence 1, Appli	Sequence 1, Appli
US-08-681-151-3	US-09-023-942A-26	US-08-944-483-63	US-08-978-404B-44	US-08-944-483-68	US-09-734-675-4	US-09-023-942A-6	US-09-387-375-7	US-08-956-267A-2	US-09-008-271A-3	US-09-023-942A-4	US-09-027-337-2	US-09-644-600-2	US-08-991-761A-13	US-09-386-653A-7	US-08-248-629A-1	US-08-451-932-1	US-08-452-260-1
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                                                                               APPLICANT: Mack, David
APPLICANT: Mack, David
APPLICANT: Mich Kurt
APPLICANT: Mison, Kurt
APPLICANT: Wilson, Kurt
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMP
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
CURRENT APPLICATION NUMBER: US/09/656,002
CURRENT FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
PRIOR FILING DATE: 2000-03-15
NUMBER OF SEQ ID NOS: 3
SOFTWARE: PATCHLIN VERSION 3.0
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97.4%; Pred. No. 5.5e-128;
iive 2; Mismatches 4;
                     Sequence 2, Application US/09656002
Patent No. 6455668
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Best Local Similarity 97.4's
Matches 228; Conservative
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ORGANISM: Homo sapiens
                                                               GENERAL INFORMATION:
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US-09-656-002-2
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61 GEWPWQASIQHPGAHVCGGSIJAPQWVLTAAHCFPRRALPABYRVRLGALRIGSTSPRTL 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              117 SLAVAKIIIIBFNPMYPKD---NDIALMKLQFPLTFSGTVRPICLPFFDEELTPATPLWI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                174 IGWGFTKQNGGKMSD--ILLQASVQVIDSTRCN----ADDAYQGE--VTEKNMCAGIPE
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                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Amino acid; OTHER INFORMATION: sequence of BOS zymogen fusion gene US-09-387-375-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 316;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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APPLICANT: Oi, Jian-shen
TITLE OF INVENTION: DNA encoding the novel human serine
TITLE OF INVENTION: Protease T
FILE REFERENCE: ORT-1032
                                                                                                       APPLICANT: Qi, Jenson
TITLE OF INVENTION: DNA Encoding the Human Serine
TITLE OF INVENTION: Protease EOS
FILE REPERENCE: ORT-1031
CURRENT APPLICATION NUMBER: US/09/387,375
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEO ID NOS: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
47.8%; Score 765; DB 4;
Best Local Similarity 51.1%; Pred. No. 3.4e-74;
Matches 164; Conservative 38; Mismatches 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/386,653A
CURRENT FILING DATE: 1999-08-31
NUMBER OF ESQ ID NOS: 11
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                                    GENERAL INFORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Andrade-Gordon, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Andrade-Gordon, Patricia
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9
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US-09-386-653A-9
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LENGTH: 315
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASISEO for Windows Version 2.0
                                                                                                                              Yue, Henry
Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 1266; DB 3;
Pred. No. 5.7e-128;
2; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Mohan-Peterson, Sheela
REGISTRATION NUMBER: 41,201
REFERENCE/DOCKET NUMBER: PF-0458 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271A
FILING DATE: 16-Jan-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: ~CURKNOWN>
FILING DATE: CINKNOWN>
ATTORNEY/AGENT INFORMATION:
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SEQUENCE DESCRIPTION: SEQ ID NO: 6
Sequence 6, Application US/09008271A
Patent No. 6203979
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
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TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 435 amino acids
TYPE: amino acid
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INFORMATION FOR SEQ ID NO: 6:
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Best Local Similarity 97.4%;
Matches 228; Conservative
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COMPUTER READABLE FORM:
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US-09-387-375-9
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; Sequence 9, Application US/09387375

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us-09-607-745-9.rai

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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 DSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSDKLGSFPSLA- 119
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                                                                                                                                        61 DSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSDKG-GSFPSLA 119
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                                                                                                                                                                                                                                                                                                                                                                        234 FEEGKKDACKGDSGGFLVCLVGQSWLQAGVISWGEGCARONRPGVIRVIAHHNWIHRII 293
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                                                                                              1 MDSKGSSQKSKLLILILLVVSNLLLCQGVVSDYKDDDDVDAALAAPFDDDDKIVGGYALEE
                                                                                                                                                                61 GEWPWQVSIQRNGSHFCGGSLIABOWVLTAAHCFRNTSETSLYOVLLGAROLVOPGPHAM
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                                                                                                                                                                                                             120 VAKIIIIEFNPMY---PKDNDIALMKLOFPLIFSGTVRPICLPFFDEELTPATPL--WII
                                                                                                                                                                                                                                                                                  GWGFTKQNGGKMSD-----ILLQASVQVIDSTRCN-----ADDAYQGE-VTEKMMCAG
                                                                                                                                                                                                                                                                                                                                                    222 IPEGGVDTCQGDSGGPLMYQSDQ-WHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNV-
                                                                      1 MDSKGSSQKSRLLLLLLVVSNLLLCQGVVSDYKDDDDVDAAALAAPFDDDDKIVGGYALDV
                                      Gaps
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                                      37;
   Length 315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46.6%; Score 746; DB 4; Length 328;
49.7%; Pred. No. 4e-72;
tive 35; Mismatches 94; Indels
                                      Indels
                                    83;
   DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/09386642
Patent No. 6420157
GRERRAL INPORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Qi, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REPERENCE: ORT-1028
CURRENT PELICANT: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 11
LENGTH: 328
47.2%; Score 754.5; DB 48.4%; Pred. No. 4.6e-73; iive 46; Mismatches 83
                                                                                                                                                                                                                                                                                                                                                                                                                         280 ----- WKAELSRHHHHH 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                        294 PKLQFQPARLGGQKSRHHHHH 315
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                  Best Local Similarity 48.43
Matches 156; Conservative
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Matches 163; Conservative
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US-09-386-642-11
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APPLICANT: Darrow, Andrew L.
APPLICANT: Qi, Jenson
APPLICANT: Adrade-Gordon, Patricia
TITLE OF INVENTION: Idenification and Characterization of the complementary
TITLE OF INVENTION: DNA encoding the novel human serine protease C-E
CURRENT APPLICATION NUMBER: US/09/386,629
CURRENT FILIG DARRE: 1999-08-31
SOFTWARE: PATCHIN ONS: 11
SOFTWARE: PATCHIN VET. 2.0
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277
                                    241 CQGDSGGPLSCPVBGLMYLTGIVSWGDACGARNRPGVYTLASSYASWIQSKVTELQPRVV 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of Artificial Sequence: C-E catalytic
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   CQGDSGGPLMYQSD-QWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIY
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APPLICANT: Qi, Jenson
APPLICANT: Qi, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
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Patent No. 6420157
GENERAL INFORMATION:
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Patent No. 6426199
GENERAL INFORMATION:
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---VRLGDHSLQNKDGP 114
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    137 -DIALMKLOFPLIFSGTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQ-AS 194
                                                                                               EQEIPVVQSIP---HPCYNSSDVBDHNHDLMLQLRDQASLGSKVKPISLA--DHCTQPG
                                                                                                                                                                                  FPSLAVAKIIIIEFNPMY-----PKDNDIALMKLQFPLTFSGTVRPICLPFFDEELTPA
                                                                                                                                                       TPLWIIGWGFTKONGGKMSDILLOASVQVIDSTRCNADDAYOGEVTEKMMCAGIPEGGVD
                                                                                                                                                                                                                                       TCQGDSGGPLMYQSDQWHVVGIVSWGYG-CGGPSTPGVYTKVSAXLNWIYNVWKABLSRH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28 VSDYKDDDDVDAAALAAPFDDDDKIVGGYALDVDSWPWQVSIQYDKQHVCGGSILDPHWV
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  PHSQPWQAALFQGQLLCGGVLVGGNWVLTAAHCKKPKYT
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42.9%; Pred. No. 1.3e-56;
ative 46; Mismatches 85;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 53, Application US/09386642
Patent No. 6420157
GANERAL INFORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Qi, Jenson
APPLICANT: Applicate-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REPERENCE: ORT-1028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/386,642
CUMRENT FILING DATE: 1999-08-31
CUMBER OF SEQ ID NOS: 60
SOFTWARE: PATCHLIN Ver. 2.0
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Patent No. 6420157
GENERAL INFORMATION:
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Best Local Similarity
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US-09-386-642-53
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US-09-386-642-12
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LENGTH: 306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           229 TCQGDSGGPLMYQSDQWHVVGIVSWGYG-CGGPSTPGVYTKVSAYLNWIYNVWKAELSRH
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                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Fusion gene; OTHER INFORMATION: with homo sapien serine protease catalytic domain US-09-386-642-14
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                                                                                                                                                                                                                                                           Length 289;
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40.3%; Score 645; DB 4; Length 288;
Best Local Similarity 47.9%; Pred. No. 2.6e-61;
Matches 146; Conservative 38; Mismatches 91; Indels
                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                    Query Match 42.0%; Score 672.5; DB 4; Best Local Similarity 48.5%; Pred. No. 2.8e-64; Matches 148; Conservative 39; Mismatches 89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Darrow, Andrew APPLICANT: Oi. Jenson APPLICANT: Oi. Jenson APPLICANT: Andrade-Goordon, Patricia PILE PEFERENCE: ORT-1028 CURRENT APPLICATION NUMBER: US/09/386,642 CURRENT FILING DATE: 1999-08-31 NUMBER OF SEC. ID NOS: 60 SOFTWARE: Patentin Ver. 2.0
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Patent No. 6420157
GENERAL INFORMATION:
           NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 14
IENGTH: 289
TYPE: PRT
ORGANIOM: Artificial Sequence
FEATURE:
CURRENT FILING DATE: 1999-08-31
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LENGTH: 288
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APPLICANT: Kazuyoshi YAMAOKA et al.
TITLE OF INVENTION: NUCLEIC ACID SEQUENCE ENCODING
TITLE OF INVENTION: TRYPSIN-LIKE ENZYME AND PROCESS FOR PRODUCING THE ENZYME
                                                                                                                                                                                                                                                                                                      61 PHSQPWQAALVMENELFCSGVLVHPQWVLSAAHCFQN----SYTIGLGLHSLEADQEPG 115
                                                                                                                                                                                                                                                                                                                                                                                                                                 118 LAVAKIIIIEFNPMYPKD---NDIALMKLQFPLIFSGTVRPICLPFFDEELTPATPLMII 174
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                                                                                                                                                                                                                                                60 VDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAGSDKL--GSFPS
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Best Local Similarity 41.8%; Pred. No. 2.5e-48;
Matches 97; Conservative 49; Mismatches 79;
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CORRESPONDERSES.
ADDRESSER: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
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MEDIUM TYPE: Diskette, 3.5 inch, 1.44
COMPUTER: IBM Compatible
OPERATING SYSTEM: NS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/508,448C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25, Application US/08508448C Patent No. 5804410 GENERAL INFORMATION:
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NAME: Warren M. Cheek, Jr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : 418 amino acids
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REFERENCE/DOCKET NUMBER
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CITY: Washington
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US-08-508-448C-25
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US-08-508-448C-25
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                                       APPLICANT: Q1, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 12
LENGTH: 319
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APPLICANT: Darrow, Andrew
APPLICANT: Q1, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.0
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Patent No. 6420157
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LENGTH: 284
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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ORGANISM: Artificial Sequence
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Matches 133; Conservative
APPLICANT: Darrow, Andrew
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US-09-386-642-54
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Score 526.5; DB 4
Pred. No. 2.5e-48;
    LUNG CANCER AND METHODS FOR
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TITLE OF INVENTION: LUNG CANCER AND METHODS; FILE REFERENCE: 210121.475C1; CURRENT APPLICATION NUMBER: US/09/370,838; CURRENT FILING DATE: 1999-08-09; EARLIER APPLICATION NUMBER: US 09/285,323; EARLIER FILING DATE: 1999-04-02; NUMBER OF SEQ ID NOS: 289; SOFTWARE: FastSEQ for Windows Version 3.0; SEQ ID NO 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
US-09-370-838-62
; Sequence 62, Application US/09370838
; Patent No. 6444425
                                                                                                                                                                                                                                                                                                      ch 32.9%;
1 Similarity 41.8%;
97; Conservative 4:
                                                                                                                                                                                                                                         ; ORGANISM: Homo sapien
US-09-370-838-83
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ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                 244 IST--TFPKLRNRVENILIHNNYKSATHENDIALVRLENSVTFTKDIHSVCLPAATQNIP 301
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                        49 DDKIVGGYALDVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWKVRAG 108
                                                                                                           109 SDKLGSFPSL--AVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEELT 166
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                                                                                                                                                                                                                                                                                  227 VDTCQGDSGGPLMYQSDQ--WHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWI 276
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APPLICANT: Reed, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Hodes, Michael J.
APPLICANT: Hodes, Michael J.
APPLICANT: Secriek. Heather
IIILE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
IIILE OF INVENTION: LOMG CANDER AND METHODS FOR THEIR USE
FILE REPRENDE: 201021, 475c1
CURRENT FILING DATE: 1999-08-09
GARLIER PILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
SOFTWARE; FASTSEQ for Windows Version 3.0
SEQ ID NO 82
LENGTH: 418
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APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadoh
APPLICANT: Secrist, Heather
IITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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41.8%; Pred. No. 2.5e-48;
ilve 49; Mismatches 79
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Patent No. 6444425
GENERAL INFORMATION:
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Patent No. 6444425
GENERAL INFORMATION:
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Best Local Similarity 41.8'
Matches 97; Conservative
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APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadoh
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE
FILE REPRENCE: 210121.475C1
CURRENT APPLICATION NUMBER: US/09/370,838
CURRENT FILING DATE: 1999-08-09
RABLIER FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
NUMBER OF SEQ ID NOS: 289
SOUTHWARE: PASESEQ for Windows Version 3.0
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227 VDTCQGDSGGPLMYQSDQ--WHVVGIVSWGYGCGGPSTPGVYTKVSAYLNNI 276

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Search completed: December 2, 2003, 02:31:10 Job time : 22.2875 secs

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US-09-888-257A-2
US-09-946-374-274
US-10-063-735-111
US-10-063-735-111
US-10-109-672-329
US-10-109-672-329
US-10-109-672-329
US-10-109-672-329
US-10-109-672-329
US-10-109-747-329
US-10-109-747-329
US-10-17-253A-274
US-10-17-253A-274
US-10-17-253A-274
US-10-17-253A-274
US-10-17-253A-274
US-10-17-253A-274
US-10-17-69-329
US-10-17-699-329
US-10-17-689-329
US-10-17-689-329
US-10-17-689-329
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Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                      US-10-175-751-329
US-10-175-754-329
US-10-176-480-329
US-10-176-489-329
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US-10-176-920-329
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Publication No. US20030166246A1
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
Hillman, Jennifer I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
  COUNTRY: USA
ZIP: 94304
 JS-10-180-719-18
                                                                                                                                                                                                                          Command line parameters:
-MODEL=frame+pol.model-DEV=xlp
-Q=/cgn2_1/USFTO_spool_p/USO9607745/runat_01122003_160940_10006/app_query.fasta_1.1038
-Q=/cgn2_1/USFTO_spool_p/USO9607745/runat_01122003_160940_10006/app_query.fasta_1.1038
-DB=Published Applications Na -QFWT=fasta_p -SUFFIX=rnpb -MINMATCH=0.1
-USOPCH=0 -LOOPEXT=0 -UNITS=blists -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40:cdi -LIST=45 -DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFWT=pto -NORM=ext -HRAPSIZE=500 -MINLEN=0
-THR_MIN=0 -ALIGN=15 -MODE=LOCAL -OUTFWT=pto -NORM=ext -HRAPSIZE=500 -MINLEN=0
-NORT=6 -LOCPU=3 -NO MAAP -LARGEQUERY -NEG SCORES=0 -WAIT -DSPBLOGK=100
-LONGLOG -DSY TIMEOUT=120 -WARN TIMEOUT=30 -THRRADS=1 -XGAPCP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOE=6 -DELEXT=7
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                                                                                     2, 2003, 02:42:05; Search time 262.68 Seconds (without alignments) 3662.446 Million cell updates/sec
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| cgn2_6/ptodata/2/pubpna/Pcr NBW Pub.seq:*
| cgn2_6/ptodata/2/pubpna/Pcr NBW Pub.seq:*
| cgn2_6/ptodata/2/pubpna/Pcr NBW Pub.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_PuB_OMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_PuB_OMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_PuB_OMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BPUBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us0_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us10_BNB PuBCOMB.seq:*
| cgn2_6/ptodata/2/pubpna/Us10_NSW PuB.seq:*
            GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                             nucleic search, using frame_plus_p2n model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2038 12 US-10-180-719-18
                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                              2190069 segs, 1647345023 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                Xgapop 10.0 , Xgapext
Ygapop 10.0 , Ygapext
Fgapop 6.0 , Fgapext
Delop 6.0 , Delext
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query
Match Length DB ID
                                                                                                                                                                                                                                                                                                              DB seq length: 0
DB seq length: 200000000
                                                                                                                                   US-09-607-745-9
1600
                                                                                      December
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Perfect score:
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                                                               protein
                                                                                                                                                              Sequence:
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Maximum |
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                                                                                      Run on:
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us-09-607-745-9.rnpb

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Sequence 2, Application US/0988257A, Publication No. US20030060612A1, GENERAL INFORMATION:
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1266.00
98.29%
97.44%
79.12%
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ORGANISM: Homo Sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Percent Similarity:
Best Local Similarity: 5
Query Match:
DB:
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US-09-888-257A-2
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                                                                                                              1403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          111 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro 130
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SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/10/180,719
FILING DATE: 25-Jun-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271
FILING DATE: 16-Jun-1998
ATTORNEY/AGENT INFORMATION:
NAMME: MOADAIN-PETERSON, Sheela
REGISTRATION NUMBER: 41,201
REFERENCE/DOCKET NUMBER: PF-0458 US
TELEPHONE: 650-855-0555
TELEPHONE: 650-855-0555
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 2038 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length:
Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
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1266.00
98.29%
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                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONE: 1337018
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Best Local Similarity:
Query Match:
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Pred. No.:
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1343 CAGGGTGACAGTGGGCCCCCTGATGTACCAATCTGACCAGTGGCATGTGGTGGGCATC 1402
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APPLICANT: Follakis Paul J.
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APPLICANT: M. Thomas D.
APPLICANT: M. THOMAS D.
APPLICANT: M. THOMAS D.
APPLICANT: M. THOMAS D.
APPLICANT: CONCEDITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF TUMOR
TITLE OF INVENTION: TREATMENT OF TUMOR
TITLE OF INVENTION: TREATMENT OF TUMOR
TITLE OF INVENTION: TREATMENT OF TUMOR
TITLE OF INVENTION NUMBER: US 60/63,540
RIOR FILING DATE: 1990-60-10
RIOR PELICATION NUMBER: US 60/09,792
RIOR FILING DATE: 1990-60-10
RIOR FILING DATE: 1990-60-10
RIOR PELICATION NUMBER: DET/US00/0534
RIOR PELICATION NUMBER: PET/US00/0534
RIOR PELICATION NUMBER: PET/US00/05328
RIOR PELICATION NUMBER: PET/US00/05666
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Conservative:
Mismatches:
Indels:
Gaps:
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Transmembrane Polypeptides and Nucleic
CTCTGGATCATTGGATGGGGCTTTACGAAGCAGAATGGGAAGATGTCTGGCATACTG 1223
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                                                                                                                                           AlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAsp
                                                                                                                                                             924 GCCCACTGCTTCAGGAAACATACCGATGTTTCAACTGGAAGGTGCGGGCAGGCTCAGAC
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                                           51 LysileValGlyGlyTyrAlaLeuAspValAspSerTrpProTrpGlnValSerIleGln
                                                                  804 CGTGTGGTGGGTGGGAGGAGGCCTCTGTGGATTCTTGGCCTTGGCAGGTCAGGATCCAG
                                                                                         TyrAsplysGlnHisValCysGlyGlySerIleLeuAspProHisTrpValLeuThrAla
                                                                                                            TACGACAACAGCACGTCTGTGGAGGAGCATCCTGGACCCCACTGGGTCCTCACGGCA
                                                                                                                                                                                            LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro
                                                                                                                                                                                                                    AAACTGGGCAGCTTCCCATCCCTGGCTGTGGCCAAGATCATCATCATTGAATTCAACCCC
                                                                                                                                                                                                                                              WetTyrProLysAspAsnAspIleAlaLeuMetLysLeuGlnPheProLeuThrPheSer
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               US-09-607-745-9 (1-292) x US-09-888-257A-2 (1-2063)
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Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
Godlowski, Paul J.
Grimaldi, Christopher J.
Hillan, Kenneth J.
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Paoni, Nicholas F.
Roy, Margaret Ann
Smith, Victoria
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Tumas, Daniel
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Eaton, Dan L.
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APPLICANT:
APPLICANT
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PRIOR FILING DATE: 1998-09-18
PRIOR APPLICATION NUMBER: 60/100849
PRIOR FLING DATE: 1998-09-17
PRIOR PELLING DATE: 1998-09-17
PRIOR PELLING DATE: 1998-09-17
PRIOR PELLING DATE: 1998-09-17
PRIOR PELLING DATE: 1998-09-18
PRIOR PELLING DATE: 1998-09-22
PRIOR PELLING DATE: 1998-09-22
PRIOR PELLING DATE: 1998-09-23
PRIOR PELLING DATE: 1998-09-24
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PRIOR FILING DATE: 1998-09-24
PRIOR APPLICATION NUMBER: 60/102207
PRIOR PILING DATE: 1998-09-29
PRIOR PELING DATE: 1998-09-29
PRIOR APPLICATION NUMBER: 60/102331
PRIOR PELING DATE: 1998-09-39
PRIOR PELING DATE: 1998-09-39
PRIOR PELING DATE: 1998-09-30
PRIOR APPLICATION NUMBER: 60/102674
PRIOR PELING DATE: 1998-09-30
PRIOR PELING DATE: 1998-10-01
PRIOR PELING DATE: 1998-10-02
PRIOR PELING DATE: 1998-10-02
PRIOR PELING DATE: 1998-10-06
PRIOR PELING DATE: 1998-10-06
PRIOR PELING DATE: 1998-10-07
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191 LeuGlnAlaSerValGlnValileAspSerThrArgCysAsnAlaAspAspAlaTyrGln 210

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APPLICANT: Exton, Dan L.

APPLICANT: Exton, Dan L.

APPLICANT: Exton, Dan L.

APPLICANT: Geritzen, Mary B.

APPLICANT: Godwski, Paul J.

APPLICANT: Godwski, Paul J.

APPLICANT: Godwski, Paul J.

APPLICANT: Grimald, Christopher J.

APPLICANT: Watanabe, Colin K.

APPLICANT: Watanabe, C
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            111 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro 130
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                                                                                                                            131 MetTyrProlysAspAspAspleAlaLeuMetLysLeuGlnPheProLeuThrPheSer
                                                                                                                                                                                                                                             151 GlyThrValArgProlleCysLeuProPhePheAspGluGluLeuThrProAlaThrPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    211 GlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAspThrCys
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US-10-063-735-111
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APPLICANT: Gurney, Austral L.
APPLICANT: Hilan, Kenneth J.
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, James
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REPRESENCE: P2830PLC54
CURRENT APPLICATION NUMBER: US/10/015,387A
CURRENT FILING DATE: 2001-12-12
Prior Application removed - See File Wrapper or Palm
SEQ ID NOS: 477
LENGTH: 2063
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Publication No. US20030135034A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Betsein, David
APPLICANT: Baton, Dan 1.
APPLICANT: Baton, Dan 1.
APPLICANT: Ferrara, Napoleone
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Clang
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
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; NUMBER OF SEQ ID NOS: 477; SEQ ID NO 274; LENGTH: 2063
; TYPE: DN A
; TYPE: DN A
; ORCANISM: Homo sapiens US-10-006-130A-274
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                                                                                                                                                                                                                       924 GCCCACTGCTTCAGGAAACATACCGATGTGTTCAACTGGAAGGTGCGGGCAGGCTCAGAC
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                                                                                         TyrAspLysGlnHisValCysGlyGlySerIleLeuAspProHisTrpValLeuThrAla
                                                                                                                                  1464 GCCTATCTCAACTGGATCTACAATGTCTGGAAGGCTGAGCTG 1505
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CURRENT FILING DATE: 2002-03-19
Prior Application removed - See File Wrapper or Palm
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APPLICANT: Paoni, Nicholas F.
TITLE OF INVENTION: Secreted and Transmembrar
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2830P1C7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Beton, Dan I.
APPLICANT: Ferrara, Napoleone
APPLICANT: Forg, Sherman
APPLICANT: Forg, Sherman
APPLICANT: Goodwakh, Paul J.
APPLICANT: Goddwad, Audrey
APPLICANT: Grimaldi, Cinistopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Hillan, Kenneth J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 274, Application US/10006130A Publication No. US20030148375A1
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US-10-006-130A-274
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1284 GGGGAAGTCACCGAGAAGATGATGTGTGCAGGCATCCCGGAAGGGGTGTGGACACCTGC 1343
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APPLICANT: Panh, James
APPLICANT: Panh, James
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
TITLE OF INVENTION: Acids Encoding the Same
CURRENT APPLICATION NUMBER: US/10/06,172A
CURRENT FILING DATE: 1998-09-01
PRIOR PILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR PILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098750
PRIOR PILING DATE: 1998-09-01
PRIOR PILING DATE: 1998-09-01
PRIOR PILING DATE: 1998-09-02
PRIOR PILING DATE: 1998-09-09
1104 GGCACAGTCAGGCCCATCTGTCTGCCTTTGATGAGGAGCTCACTCCAGCCACCCCA 1163
                                                                                                                              1164 CTCTGGATCATTGGATGGGGCTTTACGAAGCAGAATGGAGGGAAGATGTCTGACATACTG 1223
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Fong, Sherman
Gao, Weir-Qiang
Goddard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
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APPLICATION NUMBER: 60/099602
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APPLICATION NUMBER: 60/099642
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Hillan, Kenneth J
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Baton, Dan 1.
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US-10-006-172A-274
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                                                                                                                                                                                                                        APPLICANT: Zhang_Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCOING THE SAME
FILE REFERENCE: 1943OR121
CURRENT APPLICATION NUMBER: US/10/199,672
CURRENT FILING DATE: 2002-01-18
PRIOR APPLICATION NUMBER: US/10/05263
PRIOR FILING DATE: 1997-09-18
PRIOR PLING DATE: 1997-09-18
PRIOR PLING DATE: 1997-09-18
PRIOR PLING DATE: 1997-09-18
PRIOR PLING DATE: 1997-10-21
PRIOR PLING DATE: 1997-10-24
PRIOR PLING DATE: 1997-10-28
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Matches:
Conservative:
Mismatches:
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1266.00
98.29%
97.44%
                                                                                                                    Smith, Victoria
Watanabe, Colin K.
Wood, William I.
                          Godowski, Paul J.
Gurney, Austin L.
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; ORGANISM: Homo Sapien
US-10-199-672-329
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Best Local Similarity:
Query Match:
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Pred. No.:
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PRIOR FILING DATE: 1998-09-09
PRIOR PRICATION DATE: 1998-09-10
PRIOR PRICATION NUMBER: 60/099754
PRIOR PRICATION NUMBER: 60/099763
PRIOR PRICATION NUMBER: 60/099763
PRIOR PLING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-16
PRIOR PLING DATE: 1998-09-17
PRIOR PLING DATE: 1998-09-18
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09-20
PRIOR PLING DATE: 1998-09
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RAPPLICATION NUMBER: 60/102240

RETLING DATE: 1998-09-29

RAPPLICATION NUMBER: 60/102307

RETLING DATE: 1998-09-29

RAPPLICATION NUMBER: 60/102330

RAPPLICATION NUMBER: 60/102330

RAPPLICATION NUMBER: 60/102331

RETLING DATE: 1998-09-29

RAPPLICATION NUMBER: 60/102484

RETLING DATE: 1998-09-30

RAPPLICATION NUMBER: 60/102487

RETLING DATE: 1998-09-30

RAPPLICATION NUMBER: 60/102570

RAPPLICATION NUMBER: 60/102570

RAPPLICATION NUMBER: 60/102570

RAPPLICATION NUMBER: 60/102571

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RAPPLICATION NUMBER: 60/102687

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RAPPLICATION NUMBER: 60/102687 R FILING DATE: 1998-09-24
R APPLICATION NUMBER: 60/101743
R APPLICATION NUMBER: 60/101915
R APPLICATION NUMBER: 60/101915
R FILING DATE: 1998-09-24
R APPLICATION NUMBER: 60/101916
R APPLICATION NUMBER: 60/10207
R APPLICATION NUMBER: 60/102207
R APPLICATION NUMBER: 60/102207 R FILING DATE: 1998-10-02
R APPLICATION NUMBER: 60/103258
R FILING DATE: 1998-10-06
R APPLICATION NUMBER: 60/103314
R FILING DATE: 1998-10-07
R APPLICATION NUMBER: 60/103315 FILING DATE: 1998-10-07 APPLICATION NUMBER: 60/103328 FILING DATE: 1998-10-07 APPLICATION NUMBER: 60/103395 FILING DATE: 1998-10-07 APPLICATION NUMBER: 60/103396 FILING DATE: 1998-10-07 APPLICATION NUMBER: 60/103401 FILING DATE: 1998-10-07 APPLICATION NUMBER: 60/103449 FILING DATE: 1998-10-06 APPLICATION NUMBER: 60/103633 APPLICATION NUMBER: 60/103679 FILING DATE: 1998-10-08 APPLICATION NUMBER: 60/103711 FILING DATE: 1998-10-08 APPLICATION WINBER: 60/104257 FILING DATE: 1998-10-14 APPLICATION NUMBER: 60/104987 FILING DATE: 1998-10-20 APPLICATION NUMBER: 60/105000 APPLICATION NUMBER: 60/105002 FILING DATE: 1998-10-20 APPLICATION NUMBER: 60/105104 APPLICATION NUMBER: 60/105169 1998-10-08 1998-10-20 1998-10-21 FILING DATE: FILING DATE: PRIOR
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TYPE: DNA
ORGANISM: Homo Sapien
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Best Local Similarity:
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Matches:
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Indels:
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          PRIOR APPLICATION NUMBER: 60/105807
PRIOR FILING DATE: 1998-10-27
PRIOR PILING DATE: 1998-10-27
PRIOR PILING DATE: 1998-10-27
PRIOR APPLICATION NUMBER: 60/105882
PRIOR PILING DATE: 1998-10-27
PRIOR APPLICATION NUMBER: 60/106023
PRIOR FILING DATE: 1998-10-28
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; Sequence 329, Application US/10187749
; Publication No. US20030153036A1
; GENERAL INFORMATION:
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1266.00
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79.12%
PRIOR FILING DATE: 1998-10-26
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Best Local Similarity:
Query Match:
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APPLICANT: GORGNEAL, PAIL
APPLICANT: GURDEY, AUSTIN 1.
APPLICANT: GURDEY, AUSTIN 1.
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Watanabe, Colin M.
APPLICANT: Watanabe, Colin M.
APPLICANT: Chang, Zemin 1.
APPLICANT: Chang, Zemin 1.
APPLICANT: Chang, Zemin 1.
APPLICANT: Chang, Zemin 1.
APPLICANT: DATE: 2002-07-01
CURRENT APPLICATION NUMBER: US/10/187, 749
CURRENT APPLICATION NUMBER: US/10/052, 586
FRIOR APPLICATION NUMBER: 60/059263
FRIOR PILLING DATE: 1997-09-18
FRIOR PILLING DATE: 1997-09-18
FRIOR PILLING DATE: 1997-09-18
FRIOR APPLICATION NUMBER: 60/06326
FRIOR APPLICATION NUMBER: 60/06326
FRIOR APPLICATION NUMBER: 60/06321
FRIOR APPLICATION NUMBER: 60/06321
FRIOR APPLICATION NUMBER: 60/063240
FRIOR PILLING DATE: 1997-10-24
FRIOR PILLING DATE: 1997-10-28
FRIOR FILLING DATE: 199
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Desnoyers, Luc
Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Mismatches:
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                                                                                                                                                                      1266.00
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Goddard, Audrey
Godowski, Paul J
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APPLICANT: Chen, Jian
APPLICANT: Desnoyers, Luc
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul
                                                   Sapien
                                                                                                                                                                                        Percent Similarity:
Best Local Similarity:
Query Match:
                       TYPE: DNA ORGANISM: Homo
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                                                                        US-10-194-457-329
                                                                                                                          Alignment Scores:
Pred. No.:
LENGTH: 2063
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                                                         1344 caddeddacadredredeccercardracearcreaecadredeardredecardesecare 1403
                                                                                                                                                                                                                                                                                                             211 GlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAspThrCys 230
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                                                                                                                                                                                                                     LeuGlnAlaSerValGlnVallleAspSerThrArgCysAsnAlaAspAspAlaTyrGln 210
               GlyThrValArgProIleCysLeuProPhePheAspGluGluLeuThrProAlaThrPro 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Pan, James
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Matanabe.Colin K.
APPLICANT: Wood, William I.
APPLICANT: Chang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
CURRENT APPLICATION NUMBER: US/10/194,457
CURRENT APPLICATION DATE: 2002-07-11
                                                                                                                  LeuTrpileileGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAspileLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1464 GCCTATCTCAACTGGATCTACAATGTCTGGAAGGCTGAGCTG 1505
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CURRENI APPLICATION NOMBER: US/100194,45)
PRIOR APPLICATION NUMBER: 10/05286
PRIOR APPLICATION NUMBER: 10/05286
PRIOR PILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/063266
PRIOR FILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: 60/063260
PRIOR APPLICATION NUMBER: 60/06320
PRIOR APPLICATION NUMBER: 60/063120
PRIOR APPLICATION NUMBER: 60/063120
PRIOR PILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063121
PRIOR PILING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: 60/063486
PRIOR PILING DATE: 1997-10-21
PRIOR APPLICATION NUMBER: 60/063541
PRIOR APPLICATION NUMBER: 60/063541
PRIOR PILING DATE: 1997-10-28
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US-10-194-457-329
is Sequence 329, Application US/10194457
is Publication No. US20030153037A1
is GENERAL INFORMATION:
is APPLICANT: Baker, Kevin P.
is APPLICANT: Chen, Jian
is APPLICANT: Chen, Jian
is APPLICANT: Geddard, Audrey
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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Sequence 329, Application US/10196747

Publication No. US20030162250A1

GENERAL INFORMATION:

APPLICANT: Baker, Kevin P.

APPLICANT: Chen, Jian

APPLICANT: Goddard, Audrey

APPLICANT: Goddowski, Paul J.

APPLICANT: Smith, Victoria

APPLICANT: Smith, Victoria

APPLICANT: Smith, Victoria

APPLICANT: Matanabe, Colin K.

APPLICANT: Matanabe, Colin K.

APPLICANT: Amod, William I.

APPLICANT: Applicant Acids Encoping THE SAME

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC

TITLE OF INVENTION: SECRETED SECONDING THE SAME

TITLE REFERENCE: P3430RIC346

CURRENT APPLICATION NUMBER: US/10/196,747
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     284
271 AlaTyrLeuAsnTrpileTyrAsnValTrpLysAlaGluLeu
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Mismatches:
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Matches:
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ORGANISM: Homo Sapien
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Best Local Similarity:
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Pred. No.:
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                                                                                      APPLICANT: Wood, Milliam I.
APPLICANT: Wood, Milliam I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE: POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3430R1C194
CURRENT APPLICATION NUMBER: US/10/184,642
CURRENT FILLING DATE: 2002-06-27
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
LENGTH: 2063
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                    Pan, James
Smith, Victoria
Watanabe, Colin K.
Wood, William I.
Zhang, Zemin
Gurney, Austin L
                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo Sapien
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Best Local Similarity:
Query Match:
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APPLICANT:
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1104 GGCACACICAGGCCCATCTGTGTGCCCTTTTGATGAGGAGCTCACTCCAGCCACCCCA 1163
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Publication No. US20030166055A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Besnoyers, Luc
APPLICANT: Baton, Dan luc
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Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
Godowski, Paul J.
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Hillan, Kenneth J.
                98.29%
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Matches:
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-cation No. US2003016690.
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PLICATION NUMBER: US/10/173,689

CURRENT APPLICATION NUMBER: US/2002-06-17

Prior Application removed - See File Wrapper or Palm

NUMBER OF SEQ ID NOS: 612
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Publication No. US20030166104A1
GENERAL INFORMATION:
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Watanabe, Colin K.
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Pan, James
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ORGANISM: Homo Sapien
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US-10-173-689-329
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           TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE OF INVENTION: Acids Encoding the Same FILE OF INVENTION: Acids Encoding the Same FILE SPERICAL PROBLEM ACIDS ACID
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US-10-017-253A-274
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Query Match:
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LENGTH: 2063
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COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastERD for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271A
FILING DATE: 16-Jan-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: HUMAN PROTEASE MOLECULES NUMBER OF SEQUENCES: 24 CORRESPONDENCE ADDRESS:
US-08-508-448C-15
US-09-370-838-79
US-09-370-838-79
US-09-342-79
US-09-61-840-29
US-09-61-840-29
US-09-61-840-1
US-09-61-840-1
US-09-61-840-1
US-09-61-840-1
US-09-61-840-1
US-09-61-840-1
US-09-61-840-1
US-09-518-040-1
US-09-518-046-1
US-09-518-046-3
US-09-644-600-1
US-09-644-600-1
US-09-644-600-1
US-09-644-600-1
US-09-644-600-1
US-09-624-600-1
US-09-624-600-1
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STREET: 3174 Porter Dr.
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-386-653A-1
US-09-620-312D-431
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REFERENCE/DOCKET NUMBER: PF-0458 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS-09-734-675-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bandman, Olga
Hillhan, Jennifer L.
Yue, Henry
Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-008-271A-18; Sequence 18, Application US/09008271A; Sequence 18, Application US/09008271A; Patent No. 6203979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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-MODEL=frame+ p2n.model -DEV=x1p
-MODEL=frame+ p2n.model -DEV=x1p
-MODEL=frame+ p2n.model -DFUSO9607745/runat_01122003_160938_9959/app_query.fasta_1.1038
-DS=fgra_1/USPTO spool_p/USO9607745/runat_01122003_160938_9959/app_query.fasta_1.1038
-DS=fsraed_PatentE NA -QPMT=fastap -SUPFIX=rni -MINMATCH=0.1 -LGOPCGL=0
-LGOPEXT=0 -UNITS=bits -STRAT=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR SCORE=pct -THR MAR=100 -THR MINEN=0 -ALIGN=15
-MODE=LOCAL -OUTPMT=pco -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=20000000
-USER=USO9607745_GCGN 1 133 @runat_01122003_160938_9959 -NCPI=6 -ICPU=3
-NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGIGG
-DSV TIMEOUT=120 -WARN TIMEOUT=30 -THRRADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DBLOP=6 -DELEXT=7
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Sequence 2, A
Sequence 10,
Sequence 9, A
Sequence 59,
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Sequence 60,
Sequence 80,
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1 MDSKGSSQKSRLLLLLVVSN......INWIYNVWKAELSRHHHHHH 292
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11. /GGT2_6/ptodata/2/ina/5A_COMB.seq:*
12. /GGT2_6/ptodata/2/ina/5B_COMB.seq:*
31. /GGT2_6/ptodata/2/ina/6A_COMB.seq:*
41. /GGT2_6/ptodata/2/ina/6B_COMB.seq:*
42. /GGT2_6/ptodata/2/ina/PCTUG_COMB.seq:*
63. /GGT2_6/ptodata/2/ina/PCTUG_COMB.seq:*
64. /GGT2_6/ptodata/2/ina/PCTUG_COMB.seq:*
65. /GGT2_6/ptodata/2/ina/PCTUG_COMB.seq:*
                              GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                              - nucleic search, using frame_plus_p2n model
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US-09-387-375-8
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Database

111098765432

Result

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TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       191 LeuGlnAlaSerValGlnValIleAspSerThrArgCysAsnAlaAspAspAlaTyrGln 210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              251 ValSerTrpGlyTyrGlyCysGlyGlyProSerThrProGlyValTyrThrLysValSer 270
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                                                                                                                                                                                                                                    Length:
Matches:
Conservative:
Mismatches:
Indels:
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TELECOMMUNICATION INPORMATION:

TELEPHONE: 650-855-0555
TELEPAK: 650-845-4166
TELEPAK: 650-845-4166
TELEPAK: 650-845-4166
TELEPAK: 650-845-4166
TELEPAK: 650-845-4166
TENGTH: 2018 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: single
TYPE: TOPOLOGY: linear
TYPE: TERARY: COLNNOT13
TELEPAK: COLNNOT13
TERARY: COLNNOT13
TELEPAK: COLNNOT13
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Sequence 1, Application US/09656002

Sequence 1, Application US/09656002

Patent No. 645568

GENERAL INFORMATION:

APPLICANT: Mack, David

APPLICANT: Mack, David

APPLICANT: Milson, Keith

TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMPOSITIONS, AI

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS

CURRENT APPLICATION NUMBER: US/09/656,002

FILE REFERENCE: A-69108/DJB/JDD/AMS

CURRENT APPLICATION NUMBER: US 09/525,993

PRIOR FILING DATE: 2000-03-15

PRIOR FILING DATE: 2000-03-15

PRIOR FILING DATE: 2000-03-15

NUMBER OF SEQ ID NOS: 3

SOFTWARE: PatentIn version 3.0

SEQ ID NO 1

LENGTH: 2079

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253 CTCGTGTCTGAGCAGTGGTGCTGTCAGCTGCTCACTGCTTCCCCAGCGAGCACCACAAG
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613 CCTCTGATCAGCGTGTAAACTGCCTGTACAACATCGAGGCCAAGCCTGAGGAG
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Mismatches:
Indels:
Patent No. 6420157
GENERAL INFORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Q1, Jenson
APPLICANT: Q1, Jenson
APPLICANT: Andrade-Gordon, Patricia
ITILE OF INVENTION: Zymogen Activation System
FILE RFFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT PILING DATE: 1999-08-31
NUMBER: OF SEQ ID NOS: 60
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-607-745-9 (1-292) x US-09-386-642-7 (1-1169)
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                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
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Db 193 AGGGAGGGCCTGGATCGTGAGGATCGAGAAGAATGGGGACCCACCTGCGCAGGGTTCT 252 Qy 111eleuAspProHisTrpValleuThrAlaAlaHisCySPheArgLysHisThrAsp 99 253 CTGCTCACCAGCCGCTGGATGACACTGCTTCAAGGACACTCGAACAAA 312 Qy 100 ValPheAsnTrpLysValArgAlaGlySerAspLysLeuGlySerPheProSerLeuAla 119 313 CCATACCTGTTCTTGTGCTGGGGGCCTGGCAGCTGGGGAACCTGGCTTCTGG 369 Qy 120 ValAlaLysTlellelleIleGluPheAsnProMetTyrProLysAspAsn 136 370 TCCCAGAAGGTGGCTGTGGGGGCCTGGCAGCTGGGAACCTGGCTTCTGG 369 Qy 137AspTleAlaLeuMetLysLeuGlnPheProLeuThrPhSesGlyThrVal 153 bb 430 GCTGTGCAGACATGCCTTCTGGGGGCCTTCGTAATTCTTGGAGGAAGGA	192 GINALASERVAIGINVAIIIEASPSETTHTATGCYSASNALAASPASPALA 192 GINALASGTTCTATCATCTGATTGGAGGCCATCTGTACTGGCGGGGGGGG	Db 907 CGCGCTCAGGGGCCTCAGGGCCTCAGGGCTCTGGGGCTCTGGGCCCGCGCGCG
Qy 213ValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAspThr 229 Db 673 CCGCACTTTGTCCAAAAGGACTGGTGGTGGCTGGTGGAGGGGGGGCAGGACGC 732 Qy 230 CysGlnGlyAspSerGlyGlyDroLeuMetTyrGlnSerAspGlnTrpHisValVal 248 T33 TGCCAGGTGACTCTGGGGCCCACTCTGCTGGAGGGTCTTGGTACTTGTACTTGTACG 792 Qy 249 GlylleValSerTrpGlyTyrGlyCysGlyGlyDroSerThrProGlyValTyrThrLys 268 Db 733 GGCATTGTGAGCTGGAGATGCTGTGGGGCCCACTGTGGAGGTTGTACTTTCTG 852 Qy 269 ValSerAlaTyrLeuAsnTrpIleTyr	RESULT Application US/09386629 ; Sequence 2, Application US/09386629 ; Sequence 2, Application US/09386629 ; Patent No. 6426199 ; GENERAL INFORMATION: ; APPLICANT: Darrow, Andrew L. ; APPLICANT: Qi, Jenson ; TILE OF INVENTION: DAR encoding the novel human serine protease C-E ; TILE OF INVENTION: DAR encoding the novel human serine protease C-E ; TILE OF INVENTION: DAR encoding the novel human serine protease C-E ; CURRENT APPLICATION NUMBER: US/09/386,629 ; CURRENT FILING DATE: 1999-08-31 ; NUMBER OF SEQ ID NOS: 11 ; SOFTWARE: PatentIN Ver. 2.0 ; SEQ ID NO 2 LENGTH: 1166 ; TYPE: DAR ; OTHER INFORMATION: Description of Artificial Sequence ; FEATURE: ; OTHER INFORMATION: domain in a zymogen activated construct US-09-386-629-2	ignment ad. No. ore:

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OTHER INFORMATION: Description of Artificial Sequence: Fusion gene OTHER INFORMATION: with homo sapien serine protease catalytic domain -09-386-642-10	Alignment Scores: Pred. No.: 6.43e-65 Length: 1052 Score: 672.50 Matches: 148 Percent Similarity: 61.31% Conservative: 39 Best Local Similarity: 48.52% Mismatches: 89 Query Match: 42.03% Indels: 29 DB:	-607-745-9 (1-292) x US-09-386-642-10 (1-1052)	1 MethspSeriysGlySerSerdlnLysSerArgLeuLeuLeuLeuValValSerAsn 20	A]a	73 CTACTCTTGTGCCAGGGGTGTGTCTCCGACTACAAGGACGACGACGACGTGGACGGGGCC 132	41 AlaLeuAlaAlaProPheAspAspAspLysIleValGlyGlyTyrAlaLeuAsp 59	60 ValAspSerTrpProTrpGlnValSerIleGlnTyrAspLysGlnHisValCysGlyGly 79	193 AAGCACTCCCAGCCTGGCAGCCAGCCCTGTTCGAGAAGACGCGGCTACTCTGTGGGGCG 252	80 SerileLeuAspProHisTrpValLeuThrAlaAlaHisCysPheArgLys96		7HisThrAspValPheAsnTrpLysValArgAlaGlySerAspLys	#11CAUCTGGGGCAGUACAACCTUCAGAAGGAGGGGGGGGGGG	ч .	20102010111111111111111111111111111111	132 1972roysaspAsnaspilealaleudelysleudinpheProleuthr 148 13 1972ro	149 PheSerGlyThrValArgProlleCysLeuProPhePheAspGluGluLeuThrProAla 168	469 ATCACCTGGGCTGTGCGACCCTCACCTCTCCTCACGCTGTGTCACTGCTGGC 522	169 ThrProLeuTrpIlelleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAsp 188	523 ACCAGCTGCTCATTTCCGGCTGGGGCAGCACGTCCAGCCCCCAGTTACGCCTGCCT	20		9 TyrGinGlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAsp 22		ThrCysGlnGlyAspSerGlyGlyProLeuMetTyrGlnSerAspGlnTrpHisValVal 24		249 GlylleValSerTrpGlyTyrGlyCysGlyGlyProSerThrProGlyValTyrThr 267	268 LysyalSerAlaTyrLeuAsnTrplleTyrAsnValTrpLysAlaGluLeuSerArgHis 287	808 AAGTCTGCAAATATGTGGGACTGGATCAGGAGGACGATGAAGAACAATTCTAGACAT 864
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73 CTACTCTTGTGCCAGGGTGTGGTCTCCGACTACAAGGACGACGACGACGTGGACGCGCC 132
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146
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Matches:
Conservative:
Mismatches:
Indels:
                                                                                                            Sequence 9, Application US/09386642
Patent No. 6420157
GENERAL INFORMATION
APPLICANT: Darrow, Andrew
APPLICANT: Qi. Jenson
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 60
SEQ ID NO 9
SEQ ID NO 9
LENGTH: 1049
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ORGANISM: Artificial Sequence
FEATURE:
288 HisHisHisHis 292
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247 CICACGGCGCGCACTGCTTTGAAACCTATAGTGACCTTAGTGATCCCTCCGGGTGGATG 306
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541 TGGGGTACATCAAAGAGGATGAGGCACTGCCATCTCCCCACACACCCTCCAGGAAGTTCAG 600
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                                                                                                                                                                                                                                  361 TACAACCGTTACTTCGTATCGAATATCTGAGCCCTCGCTACCTGGGGAATTCACCC 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ---AlaAspAspAlaTyrGlnGly
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; OTHER INFORMATION: with homo sapien serine protease catalytic domain
US-09-386-642-8
                                                                                                                 105 ValArgAlaGlySerAspLysLeuGlySerPheProSerLeuAlaValAlaLys----
                                      LeuThrAlaAlaHisCysPheArgLysHisThrAspValPheAsn-
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Patent No. 6420157
GENERAL INFORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Qi, Jenson
APPLICANT: Qi, Jenson
APPLICANT: Alarade-Goordon, Patricia
IIIE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/386,642 COURENT FILING DATE: 1999-08-31 NUMBER OF SEQ ID NOS: 60 SOFTWARE: PATENTIN VEY: 2.0
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              ThrProLeuTrplleIleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAgp
                                                  CAGAAGTGCACCGTCTCAGGCTGGGGCACTGTCACCAGTCCCCGAGAGAATTTTCCTGAC
                                                                                                                               580 ACTÓTÓAACTGTGCAGAAGTAAAAATCTTTCCCCAGAAGAAGTGT-----GAGGATGCT
                                                                                                                                                                    TyrGlnGlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAsp
                                                                                                                                                                                                                                                 ThrCysGlnGlyAspSerGlyGlyProLeuMetTyrGlnSerAspGlnTrpHisValVal
                                                                                                                                                                                                                                                                                                                              GlyileValSerTrpGlyTyrGly---CysGlyGlyProSerThrProGlyValTyrThr
                                                                                                                                                                                                                                                                                                                                                                742 GCCATCACATCCTGGGGCTCAGACCCCTGTGGGAGGTCCGACAAACCTGGCGTCTATACC
                                                                                                                                                                                                                                                                                                                                                                                                       268 LysValSerAlaTyrLeuAsnTrpIleTyrAsnValTrpLysAlaGluLeuSerArgHis
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Conservative:
Mismatches:
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APPLICANT: Qi, Jenson
APPLICANT: Qi, Jenson
TITLE CANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT FILING DATE: 1999-08-31
NUMBER: PATENT NOS: 60
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 599
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Best Local Similarity:
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             Length:
Matches:
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Alignment Scores:
Pred. No.:
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Mismatches:
Indels:
             Sequence 60, Application US/09386642
Patent No. 6420157
GENERAL INFORMATION:
APPLICANT: Darrow, Andrew
APPLICANT: Qi, Jenson
APPLICANT: Andrade-Gordon, Patricia
TITLE OF INVENTION: Zymogen Activation System
FILE REFERENCE: ORT-1028
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT APPLICATION NUMBER: US/09/386,642
CURRENT PILING DATE: 1999-08-31
NUMBER: PATENTING DATE: 1999-08-31
NUMBER: PATENTING DATE: 1999-08-31
SOFTWARE: PATENTING VOS: 60
SOFTWARE: PATENTING VOS: 60
SOFTWARE: PATENTING VOS: 60
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Matches:
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ORGANISM: Artificial Sequence
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US-09-386-642-60
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GENERAL INFORMATION:
APPLICANT: Kazuyoshi YAMAOKA et al.
TITLE OF INVENTION: NUCLEIC ACID SEQUENCE ENCODING
TITLE OF INVENTION: TRYPSIN-LIKE ENZYME AND PROCESS FOR PRODUCING THE ENZYME
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895 GTCACCTTTACCAAAGATATCCATAGTGTGTCTCCCAGCTGCTACCCAGAATATTCCA 954
                                                             127 GluPheAsnProMetTyrProLysAspAsnAspIleAlaLeuMetLysLeuGlnPhePro
                                                                                                                                                 147 LeuThrPheSerGlyThrValArgProlleCysLeuProPhePheAspGluGluLeuThr
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                                                                                                          AACAATTATAAATCTGCAACTCATGAAAATGACATTGCACTTGTGAGACTTGAGAACAGT
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805 Fifteenth Street, N.W., #700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette, 3.5 inch, 1.44 mb COMPUTER: IBM COMPATIDLE OPERATING SYSTEM: MS-DOS SOFTWARE: Wordperfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/508,448C FILING DATE: U1y 28, 1995 CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE: FILING DATE: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-508-448C-15; Sequence 15, Application US/08508448C; Patent No. 5804410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Warren M. Cheek, Jr. REGISTRATION NUMBER: 33,367
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TELEPHONE: 202-371-8850
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth
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CITY: Washington
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COUNTRY: U.S.A.
ZIP: 20005
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745 TTCGGAAAAGCCCCGTGTGGCCAAGTTGGCGTGCCAGGTGTCTACACCCAACCTCTGCAAA
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                                       ValGlnValIleAspSerThrArgCysAsnAla----AspAspAlaTyrGlnGlyGlu
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                                                                                 GIGICGGIGGIGICIGAGGAGGICIGCAGIAAGCICIAIGACCCGCIGIACCAC-----
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APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND D.
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND D.
TITLE OF INVENTION: LUNG CANCER AND METHODS FOR FILE REFERENCE: 210121.475C1
CURRENT APPLICATION NUMBER: US 09/285,323
CURRENT FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOTE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
SOFTWARE: FRASEO for Windows Version 3.0
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526.50
62.93%
41.81%
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COGANISM: Homo sapien
US-09-370-838-80
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Mismatches:
Indels:
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                                                                                                                                     Length:
Matches:
                                                   MOLECULE TYPE: cDNA to mRNA ORIGINAL SOURCE:
                                                                                                                                   1.38e-48
526.50
62.93%
41.81%
LENGTH: 1517 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
                                                                          ORGANISM: Homo sapiens
                                                                                    ; TISSUE TYPE: tracĥea
US-08-508-448C-15
                                                                                                                                                          Percent Similarity:
Best Local Similarity:
Query Match:
                                                                                                                        Alignment Scores:
Pred. No.:
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US-09-370-838-79
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Sequence 79, Application US/09370838 Patent No. 6444425 GENERAL INFORMATION:

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167 proAlaThrProLeuTrpIleIleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMet 186
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                                                                                                   ND DIAGNOSIS OF
FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative:
Mismatches:
Indels:
APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadch
APPLICANT: Gerrist, Heather
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIATITLE OF INVENTION: LUNG CANCER AND METHODS FOR TITLE REFERENCE: 210121.475C1
CURRENT APPLICATION NUMBER: US/09/370,838
CURRENT FILING DATE: 1999-08-09
BARLIER FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-607-745-9 (1-292) x US-09-370-838-79 (1-2790)
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Matches:
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526.50
62.93*
41.81*
32.91*
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Best Local Similarity:
                                                                                                                                                                                                                                                                                                                                         LENGTH: 2790
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Alignment Scores:
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                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE FILE REPERENCE: 210121.475.05 (9) 370,838 CURRENT APPLICATION NUMBER: US 09/285,323 EARLIER FILING DATE: 1999-08-09 BARLIER FILING DATE: 1999-04-02 NUMBER OF SEQ ID NOS: 289 SOFTWARE: FastSEQ for Windows Version 3.0
                                   1248 GIGTATACTCGAGTGACAGCCTACCTTGACTGGATT 1283
265 ValTyrThrLysValSerAlaTyrLeuAsnTrpIle 276
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                            Sequence 55, Application US/09370838 Patent No. 6444425
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APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadoh
APPLICANT: Secrist, Heather
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41.38%
32.78%
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ORGANISM: Homo sapien
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Best Local Similari
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LENGTH: 1462
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Pred. No.:
                                                                                        US-09-370-838-55
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Sequence 5, Appl.
Sequence 71, Appl
Sequence 22, Appl
                    Sequence 317, App
Sequence 217, App
Sequence 634, App
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             4. US-10-097-340-317
4. US-10-11-311-217
4. US-10-11-311-217
4. US-10-11-311-217
4. US-10-264-821-634
4. US-10-264-820-1
1. US-09-76-191-3
1. US-09-76-191-3
1. US-09-76-191-3
1. US-09-76-289-1
1. US-09-76-289-1
2. US-10-065-1374-274
2. US-10-065-1378-114
2. US-10-065-1378-124
2. US-10-199-672-329
2. US-10-194-457-329
2. US-10-194-457-329
2. US-10-173-691-329
2. US-10-174-589-329
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Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Dr. CITY: Palo Alto STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-175-748-329
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US-10-175
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Publication No. US20030166246A1
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
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RESULT 1
US-10-180-719-18
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-WODEL-frame+pol.model -DEV=xlp
-Q-fcgn2 1/USFTO-spool p/USO9607745/runat 01122003_160940_10006/app_query.fasta_1.1038
-Q=fcgn2 1/USFTO_spool p/USO9607745/runat_01122003_160940_10006/app_query.fasta_1.1038
-DB=Published Applications Na -QFMT=fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=blits -START=1 -RNB=1 -MATRIX=blosum62
-LOOPCL=0 -LOOPEXT=0 -UNITS=blits -START=1 -RNB=ct -THR MAX=100
-THR MIN=0 -ALIGN=15 -MODE=LOCAL -UTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0
-MAXIEN=2000000000 -USER=USO9607745_@CGN 1 1.129 @runat 01122003 160940_10006
-LOCPU=8 -NONMAHP -LARGEQUERY -NEG SCORRES 0 -WAIT -DSPBLOCK=100
-LONGLG -DEV TIMEOUT=120 -WARN TIMEOUT=30 -THREADS=1 -XGAPCP=10 -XGAPEXT=0.5
-FGAPOP=6 -FGĀPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7
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                                                                                                                                              2, 2003, 02:42:05; Search time 391.32 Seconds (without alignments) 3662.446 Million cell updates/sec
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1: /cgn2_6/ptodata/2/pubpna/USO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
3: /cgn2_6/ptodata/2/pubpna/USO6_NEW_PUB.seq:*
4: /cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*
5: /cgn2_6/ptodata/2/pubpna/USO6_PUBCOMB.seq:*
6: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
7: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
8: /cgn2_6/ptodata/2/pubpna/USO8_NEW_PUB.seq:*
9: /cgn2_6/ptodata/2/pubpna/USO8_PUBCOMB.seq:*
10: /cgn2_6/ptodata/2/pubpna/USO9_PUBCOMB.seq:*
11: /cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
13: /cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
13: /cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
14: /cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
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16: /cgn2_6/ptodata/2/pubpna/USO0_NEW_PUB.seq:*
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16: /cgn2_6/ptodata/2/pubpna/USO0_NEW_PUB.seq:*
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16: /cgn2_6/ptodata/2/pubpna/USO0_NEW_PUB.seq:*
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17: /cgn2_6/ptodata/2/pubpna/USO0_NEW_PUB.seq:*
                    GenCore version 5.1.6
(c) 1993 - 2003 Compugen Ltd.
                                                                                                       - nucleic search, using frame_plus_p2n model
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/180,719
FILING DATE: 25-Jun-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271
FILING DATE: 16-Jan-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length:
Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                NAME: Nohan-Peterson, Sheela
REGISTRATION NUMBER: 41,201
REGISTRANCE/DOCKET NUMBER: 41,201
RELECONMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 2038 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO: 18:
US-10-180-719-18
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Query Match:
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### APPLICANT: Peter VEIBY
APPLICANT: Gordon B. MILLS
APPLICANT: Gordon B. MILLS
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Robert C. BAST, Jr.
APPLICANT: Raren LU
APPLICANT: Raren LU
APPLICANT: Raren LU
APPLICANT: Raren GLAT
TITLE OF INVENTION: Nucleic Acid Molecules and Proteins For The Identification,
TITLE OF INVENTION: Assessment, Prevention, and Therapy of Ovarian Cancer
TITLE OF INVENTION: MASSESSMENT, Prevention, and Therapy of Ovarian Cancer
TITLE OF INVENTION: NUMBER: US/10/097,340
CURRENT FILING DATE: 2001-03-14
PRIOR PELING DATE: 2001-03-14
PRIOR PELING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR APPLICATION NUMBER: 60/276,026
PRIOR PELING DATE: 2001-09-26
PRIOR PELING DATE: 2001-09-26
PRIOR FILING DATE: 2001-09-10
PRIOR FILING DATE: 2001-09-10
PRIOR PELING DATE: 2001-09-10
PRIOR PELING DATE: 2001-09-10
PRIOR PELING DATE: 2001-09-19
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Sebastian HOERSCH
Shubhangi KAMATKAR
Steve G. KOVATS
Rachel E. MEYERS
Michael MORRISEY
Peter OLANDT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: John MONAHAN
APPLICANT: Sebastian HOBE
APPLICANT: Sebastian HOBE
APPLICANT: Shubhangi KAWI
APPLICANT: Reche G. KOVAN
APPLICANT: Michael MORRIE
APPLICANT: Peter OLANDT
APPLICANT: Peter OLANDT
APPLICANT: Peter VEIBY
APPLICANT: Peter VEIBY
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Mismatches:
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Matches:
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99.79%
                       ; FEATURE:

NAME/KEY: CDS

LOCATION: (215)..(1528)

COTHER INFORMATION:

US-09-851-588-7
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                    Percent Similarity:
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Query Match:
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Oy 262 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro	302 4	322 LeuTrpIleIleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAspIleLeu	342	. 362 GlyGluValThrGluLysMetMetCysAlaGlyIlePrcGluGlyGlyValAspThrCys	Qy 382 GlnGlyAspSerGlyGlyProLeuMetTyrGlnSerAspGlnTrpHisValValGlyIle 401	61 Qy 402 61 Db 1484 463	81 523	101 583	ValArgleuSerLyaAspArgSerThrleuGlnValleuAspSerThrleuGlyAsnTrp 121	aCysargGlnMet 141 ccrgragGcagarG 703	GlyTyrSerSerLysProThrPheargalavalGlulleGlyProAspGlnAspLeuAsp 161	ValValGluIleThrGluAsnSerGinGluLeuArgMetArgAsnSerSerGlyProCys 181 ; CURRENT APPLICATION NUMBER: US/10/171,311 ; CURRENT FILING DATE: 2002-06-12 ; CURRENT FILING DATE: US/00/298,159 ; PRIOR APPLICATION NUMBER: US 60/298,159	PRIOR FILING DATE: 2001-06-13 PRIOR FILING DATE: 2001-06-13 PRIOR PILING DATE: 2001-06-1	STORT	TYTASDLYSGINHISVALCYSGIYGIYGIYGEVASSPPROHISTROVALLEUTHRAIA 241	, NAME/KBY: misc feature
		2307 434 0	o' o	LysProLe AAACCCT	le11eA] TCATAGO	euasply TGGATAA	AspG]	11uG]	erA] 	PheseralacysPheaspasnPheThrGlualaLeualaGluThral(roas 	IsnSe	LysSe		Hist.	,

1184 GGCACAGTCAGGCCCATCTGCTCTCTTTGATGAGGAGCTCACTCCAGCCACCCCA 322 LeutrpilelleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAspileLeu 1244 CTCTGGATCATTGGATGATTTGGAAGCAAATGGAGGAAATGGAGGAAGTCTCTGACATACTG 342 LeuGlnAlaSerValGlnVallleAspSerThrArgCysAsnAlaAspAspAlaTyrGln 1304 CTGCAGGCTCAGTCCAGTCATTGAGGACCAGGCAAGCAGCAAGCA	Oy 362 GlyGluValThrGluLysNetMetCysAlaGlyTleProGluGlyClyValAspThrCys 381	Qy 402 ValSerTrpGlyTyrGlyCysGlyGlyProSerThrProGlyValTyrThrLysValSer 421 Db 1484 GTTAGCTGGGGCTGCGGGGGCCCCGAGGACTATACACCAAGGTCTCA 1543 Qy 422 AlaTyrLeuAsmTrpIleTyrAsmValTrpLysAlaGluLeu 435 Db 1544 GCCTATCTAACTGCATCTACAATGTCTGGAAGGTG 1585	RESULT 5 US-10-101-510-634 ; Sequence 634, Application US/10101510 ; Publication No. US20030148295A1 ; GENERAL INFORMATION: ; APPLICANT: WANG. YIXIN	NVENTION ENCE: 15 PLICATIC LING DAT		1.01e-281 Length: 2329.00 Matches:	Ey: 99.77% Misma 99.44% Indel 12 Gaps:	US-09-607-745-2 (1-435) x US-10-10-510-634 (1-2165) QY	Qy 21. ArgileProMetGluThrPheArgLysValGlyIleProlleIleIleAlaLeuLeuSer 40	
,								m	m	m m
Alignment Scores: Pred. No.: Pred. No.: 2337 Score: 2307 Matches: Percent Similarity: 100.00% Mismatches: 99.79% Indels: 0 Usery Match: 14 Gaps: US-09-607-745-2 (1-435) x US-10-171-311-217 (1-2307)	Qy 2 AspProAspSerAspGlnProLeuAsnSerLeuAspValLySProLeuArgLysProArg 21 Db 284 GATCCTGACAGTGATCAACCTCTGAACAGCCTCGATGTCAAACCCTGGGCAAACCCGT 343 Qy 22 IleProMetGluThrPheArgLysValGlyIleProIleIleIleAlaLeuGserLeu 41 Db 344 ATCCCCATGGAGACCTTCAGAAAGGTGGGGAATCCCCATCATCATACACTGAACCTG 403	aSerileileilevalvalvalueullebysvalileLeukspLysTyrTyrPheLeu aAdrarcarcarcarcarcarcaadgrafrendgaraaracracracrecre scaracracrationedraphericrecreansproprecrecrecrecrecrecrecrecrecrecrecrecrecr	Oy 82 ProLeuGlyGluAspGluGluHisCysValLysSerPheProGluGlyProAlaValAla 101	Db 584 GTCCGCCTCTCCAAGGACCGATCCACACTGCAGGTGCTGGACTCGGCCACAGGGAACTGG 643 Qy 122 PheSerAlaCysPheAspAsnPheThrGluAlaLeuAlaGluThrAlaCysArgGlnMet 141 Db 644 TTCTCTGCCTGTTTCGACAACTTCACAGAAGCTCTCGCTGAGACGCTGTAGGCAGATG 703	Oy 142 GlyTyrSerSerIsProThrPheArgAlaValGluIleGlyProAspGlnAspLeuAsp 161	Db 764 GTIGITIGAAATCACAGAAACAGCCAGGGGCTTCGCGGAACTCAAGTGGGCCCTGT 823 Oy 182 LeuSerGlySerLeuValSerLeuHisCysLeuAlaCysGlyLysSerLeuLysThrPro 201 Db 824 CTCTCAGGCTCCCTGGTCTCCCTGCATGCTTGCCTGGGAAGAGCCTGAAAGACCCCC 883	legin rccag	222 1yraspryscinisvalcysciyciyserilebouaspyronisirovalbeni 944 TACGACAACAGCACCTCTGGGGGGGGGCTAICTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT	1004 GCCCACTGCTTCAGGAAACATACCGATGTTCAACTGGAAGGTGCGGGCAGGCTCAGCACACCACTACCAGAC 262 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro	DD 1064 AAACTGGGCAGCTTCCCATCCCTGGCTGTGGCCAACGATCATCATCATGATTCAACTCC 1123 OY 282 MetTyrProLysAspAsnAspIleAlaLeuMetLysLeuGlnPheProLeuThrPheSer 301

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RESULT 6
US-09-851-588-5
Sequence 5, Application US/09851588
Sequence 5, Application US/09851588
Sequence 5, Application US/09851588
Sequence 5, Application US2002042667A1
GENERAL INFORMATION:
APPLICANT: Mack, David
APPLICANT: Wilson, Keith E.
TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMPOSITIONS, AI TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AI TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS
CURRENT APPLICATION NUMBER: US/09/851,588
CURRENT PILING DATE: 2000-09-24
SRIOR FILING DATE: 2000-08-17
PRIOR FILING DATE: 2000-08-17
PRIOR FILING DATE: 2000-09-06
NUMBER OF SEQ ID NOS: 9
SOFTWARE PATENTIN Version 3.1
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// ORGANISM: Homo sapiens
US-09-851-588-5
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Best Local Similarity:
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LENGTH: 2079
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                            CysProLeuGlyGluAspGluGluHisCysValLysSerPheProGluGlyProAlaVal
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ORGANISM: Homo sapien
FEATURE:
NAME/KEY: CDS
LOCATION: (251)...(1522)
OTHER INFORMATION: Nucleotide sequence encoding tra
OTHER INFORMATION: protease, serine 4 (TMPRSS4)
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank NM016425
DATABASE ENTRY DATE: 2000-11-06
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Mismatches:
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Matches:
   PRIOR APPLICATION NUMBER: 09/657,968
PRIOR FILING DATE: 2000-02-08
NUMBER OF SEQ ID NOS: 72
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 71
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APPLICANT: Edwin L. Madison
APPLICANT: Edwin L. Ong
APPLICANT: Edwin L. Ong
APPLICANT: Edwin Corvas International, Inc.
TITLE OF INVENTION: TRANSMEMBRANE SERINE PROTEASES, THE ENCODED PROTEINS AND
TITLE OF INVENTION: METHODS BASED THEREON
TITLE OF INVENTION: METHODS BASED THEREON
TITLE OF INVENTION: METHODS BASED THEREON
TITLE OF INVENTION: MOMBER: US/09/776,191
CURRENT FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/213,124
PRIOR APPLICATION NUMBER: 60/213, 840
PRIOR PRILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/179,982
PRIOR FILING DATE: 2000-06-22
PRIOR FILING DATE: 2000-02-03
PRIOR FILING DATE: 2000-02-18
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Publication No. US20030119168A1
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; TYPE: DNA ; CRGANISM: Homo sapiens ; FEATURE: ; OTHER INFORMATION: CJA8 CDNA	20-22	red. No.: core: ercent Similarity:	Local Similarity: 99.77* Mismatches: Match: 99.23\$ Indels: 14 Gaps:	US-09-607-745-2 (1-435) x US-10-264-820-22 (1-2079) Qy	217 GAICCTGACAGTGAICAACTCTGAACGCCTCGAIGTCAAACGCCTGCGGTGTCAAACGCCGGTCCCGTCCC	277 ATCCCCATGGAGACCTTCAGAAAGTGTGGGGATCCCCATCATCATAGCACTACTGAGCCT 33 41 uAlaSerileileileileValValvalieuileLysValileLeuAspLysTyrTyrPheLe 61	53. GGCGGGGTGTTGTGGGTTGTCGTGGGGTGTTCTGGGATGATACTTCCT 5 61 uCygG1yG1nProLeuHisPhe1leProArg1ysG1nLeuAspCy 8	397 CTGCGGGCAGCTCTCCACTTCATCCCGAGGAGCAGCTGTGTGACGGAGAGCTGGACTG 45 81 sProleuglyGluaspGludluHisCysVallysSerPheProGluglyProAlaVala1 10	457 TCCCTTGGGGGGGGGGGGGCACTGTGTCAAGAGCTTCCCCGAAGGGCCTGCAGTGGC 51 101 aValargLeuSerLysAspargSerThrLeuGlnValLeuAspSerAlaThrGlyAsnTr 12	DD 517 AGICCUCTCTCCAAGGACCGATCCACTGCAGGTGCTGGACTCGGCCACAGGGAACTG 578 QY 121 pPheSerAlaCysPheAspAsnPheThrGlualateualaGluThrAlaCysArgGlnMe 141	Oy 141 tGlyTyrSerSerLysProThrPheArgAlaValGluIleGlyProAspGlnAspLeuAs 161	Oy 161 pvalValGlulleThrGluAsnSerGlnGluLeuArgMetArgAsnSerSerGlyProCy 181	Oy 181 sLeuSerGlySerLeuValSerLeuHisCysLeuAlaCysGlyLysSerLeuLysThrPr 201	201 OARGYALOVALGIYGIUGIUALASEVALASSESTITEPROTITEGINVALSESTILGG 22 1	Qy 221 nTyraspLysGlnHisValCysGlyGlySerIleLeuaspProHisTrpValLeuThral 241 Db 877 GTACGACACACACACTCTGGAGGAGCATCCTGGACCCCCACTGGACCTCTCACGGC 936	Qy 241 aAlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAs 261	261 pLysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleGluPheAsnPr 28
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2 ASPPROASPSERASpGInProLeuAsnSerLeuAspVallysProLeuArgLysProArg 21 17 GATCCTGACATGATCACTCTGAACAGCCTCGATGTCAAACCCCTGGCGAAACCCTGT 22 IleProMetGluthrPheArglys-ValGlyIleProlleIleIleAlaLeuLeuSerLe 41 17 ATCCCCATGGGACTCTCAGAAAGTGTGGGATCCCCATCATAGCACTACTGAGCCT 33 14 uAlaSerIleIleIleValValValValLeuIleLysValIleLeuAspLysTyrTyrPheLe 61 18	161 pValValGiulleThrGluAsnSerGlnGluLeuArgAenSerSerGlyProcy 181 697 TGTTGTAGAAATGAGAAAACAGGGGGGTCTGCAGGAAATCAAGTGGGCCCTG 756 181 SLeuSerGlySerLeuValSerLeuHisCysLeuAlaCysGlyLysSerLeuLysThrPr 201 757 TCTCTCAGGCTCCCTGCACTGTCTTGCCTTGGGAAGTCCTGAGGCCCTG 716 201 OArgValValGlyGluGluAlaSerValAspSerTTpProTrpGlnValSerIleGl 221 817 CCGTGTGGTGGGTGGGAGGAGCTCTGTGGATTCTTGGCTTGGCAGTCAGCATCCA 876 221 ITYTASDLySGlnHisValCysGlyGlyGlySerIleLeuAspProTrpGlnValLeuThrAl 241 817 CCGTGTGGTGGGGGGGGGGGCTCTGTGGATTCTTGGCTTGGCAGTCCAGGATCCACGGGTCAGCATCCAGGGCTCAGGGCTCTGGGATTCTTGGCTTGGCAGTCCAGGGCTCCAGGGGCAGGCTCTGGGATTCTTGGCTTGGCATTGGATCCTCAGGGCTCAGGGGTCAGGTCAGGGTCAGGGTCAGGGTCAGGATCTTGGATTCTTGGCTTGGGATCCTCAGGGCTCAGGGGGGGG	321 oLeuTrpIleIleGlyTrpGlyPheThrLysGlnAsnGlyGlyLysMetSerAspIleLe 341 1177 ACTCTGGATCATTGGATGGGGCTTTACGAAGCAGAATGGAGGAAGATGTCTGACATACT 1236 341 uLeuGlnAlaSerValGlnValIleAspSerThrArgCysAsnAlaAspAspAlaTyrGl 361 1237 GCTGCAGGCTCAGGTCATTGACAGCACACGGTGCAATGCAGAGCGTACCA 1296
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Oy 361 nGlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAspThrCy 381		
1297 GGGGAAAGTCACCGAAGATGATGTGTTGTTGTTGTTGTTGTTGTTGTTGTTGT	g &	590 GCGAGIATCATCATTGTGGGTIGTCCTCATCCAGGTGGTTGTGGGATACTACTTTCTC 62 CygGlyGlaPtroledHisPheTleProArgLygGlaLecygAAspGlyGluLeuAspCys 62 CygGlyGluProledHisPheTleProArgLygGlaLecygAAspGlyGluLeuAspCys 63 CygGlyGluProledHisPheTleProArgLygGlaLecygAAspGlyGluLeuAspCys 63 CygGlyGluProledHisPheTleProArgLygGlaLecygAAspGlyGluLeuAspCys 64 CygGlyGluProledHisPheTleProArgLygGlaLecygAAspGlyGluLeuAspCys 65 CygGlyGluProledHisPheTleProArgClygGlaLecygAAspGlyGluCeuAspCycluCcus 65 CygGlyGluProledHisPheTleProArgClygGluProledHisPheTle
UN 381 SCHOLYAPPSETVYSTYPTOLENBETVYSTINSETASDEINTINTSVALAULISTA 401 Db 1357 CCAGGGTGACAGTGGTGGGCCCCTGATGTACCAATCTGACCAGTGGCATTGGGTGGCAT 1416	අග	- HG .
Qy 401 eValSerTrpGlyTyrGlyClyGlyClyProSerThrProGlyValTyrThrLysValSe 421 Db 1417 CGTTAGCTGGGGGCTGCGGGGCCCCGAGCACCCCAGGAGTATACACCAAGGTCTC 1476	Qy Dp	82 ProfeudlyGluAspGludluHisCysYalLysSerPheProGludlyProAlaValAla 1
421 ralaTyrLeuAsnTrplleTyrAsnValTrpLysAlaGluLeu 435	. da	102 Valargleuserlysäspargserthrleuglnvalleukspseralathrelyäsntrp 1
SULT 10 -09-776-191-3	Oy.	122 PheSerAlaCysPheAspAsnPheThrGlualaLeuAlaGluThrAlaCysArgGlnMet 1
Publication No. US20030119168A1 GENERAL INFORMATION: APPLICANT: Edwin L. Madison	Oy Db	142 GlyTyrSerSerLysProThrPheArgAlaValGluIleGlyProAspGlnAspLeuAsp 1
SNIGOOD	Qy	162 ValValGluIleThrGluAsnSerGlnGluLeuArgMetArgAsnSerSerGlyProCys
TITLE OF INVENTION: ACADOMICATION OF THEREON FILE REFERENCE: 24745-1607 CURRENT APPLICATION NUMBER: US/09/776,191	QY DP	182 LeuSerGlySerLeuValSerLeuHisCysLeuAlaCysGlyLysSerLeuLysThrPro 3
CORRENT FILING DAIE: 2001-02-02 PRIOR PELING DAIE: 60/213,124 PRIOR FILING DAIE: 2000-06-22 PRIOR APPLICATION NUMBER: 60/234,840	O.V.	202 ArgvalValGlyGlyGluGluAlaSerValAspSerTrpProTrpGlnValSerTleGln 2
; FRIOR FILING DATE: 2000-06-22 ; PRIOR FILING DATE: 2000-02-03 ; PRIOR APPLICATION NUMBER: 60/183,542	Qy	222 TyraspLysGlnHisValCysGlyGlySerIleLeuaspProHisTrpValLeuThrala 2
FALOR FILLING DATE: 2000-02-18 FRIOR FILLING DATE: 2000-02-08 NUMBER OF SEQ ID NOS: 72 NUMBER OF SEQ ID NOS: 72	Qy	242 AlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAsp 2
SEQ ID NO 3 LENGTH: 2 TYPE: DNA	Qy	262 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsnPro 3
) OKCANISM: Homo Sapien	λ dg	282 MetTyrProLysaspAsnAspIleAlaLeuMetLysLeuGlnPheProLeuThrPheSer 3
orocease (MTSP3) protein	Qy Db	302 GlyThrValargProlleCysLeuProPhePheAspGluGluLeuThrProAlaThrPro 3
Alignment Scores: Pred. No.: 1.79e-280 Length: 2137 Score: 2319.00 Matches: 432 Percent Similarity: 99.54 Conservative: 0	\$ Q	
99.02% Intellations: 11 Gaps: x US-09-776-191-3 (1-2)37)	\text{\tint{\text{\tin}\text{\ti}\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tin}\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\tin}\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\text{\ti}\text{\text{\text{\text{\texi}\text{\text{\texi}\text{\text{\text{\tex{\text{\text{\text{\text{\text{\texi}\text{\texi}\text{\text{\text{\text{\text{\text{\text{\texi}\text{\text{\texi}\text{\t	342 LeuGlnAlaSerValGlnVall1eAspSerThrArgCysAsnAlaAspAsaPalaTyrGln 3
AspProAspSerAspGlnProLeuAsnS 	Qy	362 GlyGluValThrGluLysMetMetCysAlaGlyIleProGluGlyGlyValAspThrCys 3 [
22 IleproMetGluThrPheArgLySValGlyIleProllellelleAlaLeuLeuSerLeu 4	Š a	382 GlnGlyAspSerGlyGlyProLeuMetTyrGlnSerAspGlnTrpHisValValGlyIle 4
61	م 	402 ValSerTrpGlyTyrGlyCysGlyGlyBroSerThrProGlyValTyrThrLysValSer 4

 361 1349

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161 683 181 743 803 221 863 241

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1284 GGGGAAGTCACCGAGAAGATGATGTGTGCGGCAGCCTCCCGGAAGGGGGTTGGCACGCTGC 1343
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                                                   22 IleProMetGluThrPheArgLysValGlyIleProIleIleIleAlaLeuLeuSerLeu
                                                                                                      279 ATCCCCATGGAGACCTTCAGAAAGGIGGGGATCCCCATCATCATAGCACTGAGCCTG
                                                                                                                                                                                                                                                                                339 GCGAGTATCATCATGTGTTGTCCTCATCAGGTGATTCTGGATAAATACTACTTCCTC
                                                                                                                                                                                                                                                                                                                                                    62 CysGlyGlnProleuHisPhelleProArglysGlnLeuCysAspGlyGluLeuAspCys
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               162 ValValGluIleThrGluAsnSerGlnGluLeuArgMetArgAsnSerSerGlyProCys
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APPLICANT: Godowaki, Paul J
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PRIOR APPLICANTON NUMBER: US 60/063,540
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICANTON NUMBER: US 60/099,792
PRIOR FILING DATE: 1998-09-10
PRIOR PLING DATE: 1998-09-10
PRIOR APPLICANTON NUMBER: US 60/039,792
PRIOR APPLICANTON NUMBER: US 60/039,792
PRIOR PLING DATE: 1998-09-10
PRIOR APPLICANTON NUMBER: US 60/039,792
PRIOR APPLICANTON NUMBER: US 60/039,732
PRIOR APPLICANTON NUMBER: PCT/USSO(04342)
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PRIOR APPLICANTON NUMBER: PCT/USSO(0439)
PRIOR PILING DATE: 1000-03-03
PRIOR APPLICANTON NUMBER: PCT/USSO(0439)
PRIOR PILING DATE: 2000-03-03
PRIOR APPLICANTON NUMBER: PCT/USSO(05841)
PRIOR APPLICANTON NUMBER: PCT/USSO(05801)
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Mismatches:
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Matches:
                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/0988257A
Publication No. US20030060612A1
GENERAL INFORMATION:
APPLICANT: Goddard, Audrey
APPLICANT: Godwski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
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ORGANISM: Homo Sapien
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Best Local Similarity:
Query Match:
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THE APPLICATION NUMBER: 00/100310

RAPPLICATION NUMBER: 60/100930

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RAPPLICATION NUMBER: 60/101014

RAPPLICATION NUMBER: 60/101068

RETLING DATE: 1998-09-18

RETLING DATE: 1998-09-18

RAPPLICATION NUMBER: 60/101071

RAPPLICATION NUMBER: 60/101071

RAPPLICATION NUMBER: 60/101279

RETLING DATE: 1998-09-18

RETLING DATE: 1998-09-18

RETLING DATE: 1998-09-18

RETLING DATE: 1998-09-18
                                          R FILING DATE: 1998-09-10
R APPLICATION NUMBER: 60/099808
R FILING DATE: 1998-09-10
R FILING DATE: 1998-09-10
R FILING DATE: 1998-09-10
R APPLICATION NUMBER: 60/099815
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DR APPLICATION NUMBER: 60/099816

DR APPLICATION NUMBER: 60/100385

DR FILING DATE: 1998-09-10

R PLING DATE: 1998-09-10

DR FILING DATE: 1998-09-15

DR FILING DATE: 1998-09-15

DR FILING DATE: 1998-09-15

DR APPLICATION NUMBER: 60/100390

R FILING DATE: 1998-09-16

DR APPLICATION NUMBER: 60/10064

DR APPLICATION NUMBER: 60/100661

DR APPLICATION NUMBER: 60/100661

DR APPLICATION NUMBER: 60/100661

DR FILING DATE: 1998-09-16

DR PILING DATE: 1998-09-16

DR APPLICATION NUMBER: 60/100664

DR APPLICATION NUMBER: 60/100664

DR APPLICATION NUMBER: 60/100664

DR APPLICATION NUMBER: 60/100664

DR APPLICATION NUMBER: 60/100683

R FILING DATE: 1998-09-17

DR APPLICATION NUMBER: 60/100710

DR FILING DATE: 1998-09-17

DR APPLICATION NUMBER: 60/100848
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R APPLICATION NUMBER: 60/101472

R APPLICATION NUMBER: 60/101474

R FILING DATE: 1998-09-23

R APPLICATION NUMBER: 60/101474

R FILING DATE: 1998-09-23

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R FILING DATE: 1998-09-23

R APPLICATION NUMBER: 60/101476

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APPLICATION NUMBER: 60/100849
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APPLICATION NUMBER: 60/100919
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APPLICATION NUMBER: 60/101741
FILING DATE: 1998-09-24
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APPLICATION NUMBER: 60/101915
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APPLICATION NUMBER: 60/101916
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                                          1344 CAGGTGACAGTGGGCCCCTGATGTACCAATCTGACCAGTGGCATGTGGTGGCCATC 1403
                                                                                                                                                                                       1404 GITAGCIGGGGCIATGGCIGCGGGGCCCGAGCACCCCAGGAGTATACACCAAGGICICA 1463
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APPLICANT: Tunas, Daniel
APPLICANT: Tunas, Daniel
APPLICANT: Tunas, Daniel
APPLICANT: Watanabe, Colin, W.
APPLICANT: Watanabe, Colin, W.
APPLICANT: Watanabe, Colin, W.
APPLICANT: Watanabe, Colin, W.
APPLICANT: Wood, Williama I.
TITLE OF INVENTION: Acids Encoding the Same
TITLE OF INVENTION: Acids Encoding the Same
TITLE OF INVENTION: Acids Encoding the Same
FILE REPERRING: P2830PLG.
CURRENT APPLICATION NUMBER: 60/099716
FRIOR APPLICATION NUMBER: 60/099716
FRIOR PLING DATE: 1998-09-01
FRIOR PLING DATE: 1998-09-01
FRIOR PLING DATE: 1998-09-01
FRIOR APPLICATION NUMBER: 60/09863
FRIOR PLING DATE: 1998-09-00
FRIOR FILING DATE: 1998-09-00
FRIOR APPLICATION NUMBER: 60/099642
GlnGlyAspSerGlyGlyProLeuMetTyrdlnSerAspGlnTrpHisValValGly1le 401
                                                                                                                                            421
                                                                                                                               402 ValSerTrpGlyTyrGlyCysGlyGlyProSerThrProGlyValTyrThrLysValSer
                                                                                                                                                                                                                                                                                                                       1464 GCCTATCTCAACTGGATCTACAATGTCTGGAAGGCTGAGCTG 1505
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Fong, Sherman
Gao, Weil-Oiang
Goddwrd, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
Gurney, Austin L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 274, Application US/09946374; Publication No. US20030073129A1; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: 60/099741
PRIOR FILING DATE: 1998-09-10
PRIOR APPLICATION NUMBER: 60/099754
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APPLICATION NUMBER: 60/099763
FILING DATE: 1998-09-10
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Paoni, Nicholas F.
Roy, Margaret Ann
Smith, Victoria
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan L.
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APPLICANT:
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	Length:	Matches: Conservative:
1998-09-24 NUMBER: 60/102207 1998-09-29 NUMBER: 60/102240 1998-09-29 NUMBER: 60/103307 1998-09-29 NUMBER: 60/102307 1998-09-29 NUMBER: 60/102331 1998-09-29 NUMBER: 60/102331 1998-09-29 NUMBER: 60/102484	NUMBER: 60 1998-09-3 1998-09-3 1998-09-3 1998-10-0	₩ 0
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Š	102	ValArgLeuSe	r. L.	erThrLeuGlnVa	ThrGlyAsnTr
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λο ·	O	PheSerAlaCys	44	ThrGluAlaLe	CysArgGlnMet 14
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QC	684	Gricrica	CACAGAA	CAGGAGCTTCC	TCAAGTGGGCCCTG
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q	864	TACGACAAAC			
λo	242	AlaHisCys	sPheArglysH	ChrAspValPheAsnT	erAs
qq	924	CCACTG	CTTCAGGAAAC	TCAACT	SGTGCGGGCAGGCT
δλ	(i)	LysLeuGly	ySerPheProSer	erLeuAlaValAlaLysIleIle	
d d	984	AAACTGGG	Ā	Greecea	arcarreaarreaacce
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8 1	322	Leufrpile	H-1	eGlyTrpGlyPheThrLysGlnAsnGly	GlyLysMetSerAspileLeu 34
QQ	1164	CTGGAT	AI	scritacgaagcagaatgg	śdakakarcickackieci

339 GCGAGTATCATCATCATCTAGGTGATTCTGGATAAATACTACTCCTC 62 CysGlyGlnProLeuHisPhelleProArgLysGlnLeuCysAspGlyGluLeuAspCys 11	639 GGCTACAGG- 162 ValvalGluI 684 GTTGTTGAAA 182 LeuSerGLYS 744 CTCTCAGGCT 202 ArgValValQ 804 CGTGTGGTGG 222 TYTASPLYSG 222 TYTASPLYSG 864 TACGACAAAC	242 AlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAsp 26 242 GCCGCTGTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTT	322 322 1164 342 1224 362 1284 1344 1344
4 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	8 8 8 8 8 8 8 8	6 6 6 6 6 6 6	8 6 8 6 8 6 8 6 8 6
QY 342 LeuGlnAlaSerValGlnValI1eAspSerThrArgCysAsnAlaAspAspAlaTyrGln 361 Db 1224 CTGCAGGCGTCAGTCCAGGTCAATGCAGGCGAATGCGAGCGTAACCAGGTCACCAGGTAACCAGGAAGTCAACCAGAAGTCAACCAGAAGTCAACCAGAAGTCAACCAGAAGTCAACCAGAAGTCAACCAGAAGTCAACCAGGAAGTCAACCAGGAAGTCAACCAGAACTCAACCAAGAACATCAAACAATGAACCAAGAACATCAAACAAA	RESULT 13 US-10-015-387A-274 US-10-015-387A-274 Sequence 274, Application US/10015387A Publication No. US20030135034A1 GENERAL INFORMATION: APPLICANT: Befser, Kevin P. APPLICANT: Betstein, David APPLICANT: Ferrara, Napoleone APPLICANT: Ferrara, Napoleone APPLICANT: Fong, Sherman APPLICANT: Goddard, Audrey APPLICANT: Goddard, Audrey APPLICANT: Goddard, Paul J. APPLICANT: Goddwski, Paul J. APPLICANT: Girmaldi, Christopher J. APPLICANT: Girmaldi, Christopher J. APPLICANT: Girmaldi, Christopher J. APPLICANT: Hillan, Kenneth J.	HEPLICANT: Pan, James APPLICANT: Paoni, Nicholas F. ITILE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic ITILE OF INVENTION: Acids Encoding the Same CURRENT APPLICATION NUMBER: US/10/015,387A ICURRENT PILING DATE: 2001-12-12 INVERENT PILING DATE: 2001-12-12 INVERENT PILING DATE: 2001-12-12 INVERENT OS SEQ ID NOS: 477 SEQ ID NO 274 ILENGTH: 2063 ITENGTH: 2063 ITENGTH: 2063 ITENGTH: 2063 ITENGTH: 2063 INVERENT HOMO Sapiens US-10-015-387A-274	Alignment Scores: Pred. No.: 2297.50 Watches: 429 Score: 50 Best Local Similarity: 98.85* Mismatches: 0 Secore: 10

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                                                                                                              182 LeuSerGlySerLeuValSerLeuHisCysLeuAlaCysGlyLysSerLeuLysThrPro
                                                                                                                                                                                                                                                                                                                    222 TyrAspLysGlnHisValCysGlyGlySerIleLeuAspProHisTrpValLeuThrAla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 282 MetTyrProLysAspAsnAspIleAlaLeuMetLysLeuGlnPheProLeuThrPheSer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1104 GGCACAGTCAGGCCCATCTGTCTTCTTTTGATGAGGAGCTCACTCCAGCCACCCCA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ValValGluIleThrGluAsnSerGlnGluLeuArgMetArgAsnSerSerGlyProCys
                                                     684 GTTGTTGAAATCACAGAAAACAGCCAGGAGCTTCGCATGCGGAACTCAAGTGGGCCCTGT
                                                                                                                                                                                                                      ArgValValGlyGlyGluGluAlaSerValAspSerTrpProTrpGlnValSerIleGln
                                                                                                                                                                                                                                                                 804 CGTGTGGTGGGTGGGGGGGGCTCTGTGGATTCTTGGCCTTGGCAGGTCAGCATCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                        242 AlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAsp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                262 LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleGluPheAsnPro
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Godowski, Paul J.
Grimaldi, Christopher J.
Gurney, Austin L.
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Eaton, Dan 1.
Ferrara, Napoleone
Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
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APPLICANT: Botstein, David
APPLICANT: Beton, Dan 1.
APPLICANT: Eaton, Dan 1.
APPLICANT: Ferrara, Napoleon
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
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US-10-006-130A-274
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                                                                                                                                              Sequence 111, Application US/10063735

Sequence 111, Application No. US2003013882A1

GENERAL INFORMATION:

APPLICANT: Eaton, Dan L.

APPLICANT: Eaton, Dan L.

APPLICANT: Geritzen, Mary E.

APPLICANT: Godowski, Paul J.

APPLICANT: Grimaldi, Christopher J.

APPLICANT: Watanabe, Colin K.

APPLICANT: Watanabe, Colin K.

APPLICANT: Watanabe, Colin K.

TITLE OF INVENTION: ACTOR ENCODING THE SAME

FILE REFERENCE: P3230R1C1

CURRENT APPLICATION NUMBER: US/10/063,735

CURRENT APPLICATION NUMBER: US/10/063,735

CURRENT APPLICATION NUMBER: US/10/063,735

NUMBER OF SEQ ID NOS: 170

SEQ ID NO 11

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; ORGANISM: Homo Sapien
US-10-063-735-111
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Best Local Similarity:
                                                                                                        RESULT 14
US-10-063-735-111
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APPLICANT: Hillan, Kenneth J.
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, James
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P283 OPP.C7
CURRENT APPLICATION NUMBER: US/10/006,130A
CURRENT FILING DATE: 2002-03-19
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 477
SEQ ID NO 274
LENGTH: 2063
TYPE: DAR
CREANISM: Homo sapiens
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242 AlaHisCysPheArgLysHisThrAspValPheAsnTrpLysValArgAlaGlySerAsp
                                                                                                           LysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleGluPheAsnPro
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Minimum DB Maximum DB

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COMPUTER: IBM COMPALIALE
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
US-09-820-002-1

US-09-300-846-1

US-09-300-846-1

US-09-300-838-19

US-09-370-838-79

US-09-370-838-79

US-09-370-838-79

US-09-370-838-79

US-08-200-900h-1

US-08-807-151-2

US-08-10616-1

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US-09-1051-2

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JS-08-358-928-18
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REGISTRATION NUMBER: 41,201
REFERENCE/DOCKET NUMBER: PF-0458 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271A
FILING DATE: 16-Jan-1998
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ALIGNMENTS
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FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 18, Application US/09008271A; Patent No. 6203979; GENERAL INFORMATION:
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Hillman, Jennifer
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Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
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COMPUTER READABLE FORM:
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-LODEEXT=0 -UNITS=bits -GTRRT=1 -PNATRIX-blosum62 -TRANS=human40.cdi
-LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_NIN=0 -ALIGN=15
-MODEL=COCLL_OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USRES_LOCCL_OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
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-FGARDEXT=7 -YGARDOP=10 -YGARDEXT=0.5 -DBLOR=6 -DBELEXT=7
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Sequence 3, Appli
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Sequence 188, App
Sequence 188, App
Sequence 3, Appli
                                                                                                                                               2003, 02:30:15; Search time 83.7689 Seconds (without alignments) 2292.040 Million cell updates/sec
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/cgn2_6/pbcdata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/pbcdata/2/ina/Packfiles1.seq:*
                GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-861-966-188
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US-09-342-749-29
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US-09-518-749-1
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Perfect score:
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Jatabase

Result

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Sequence 1, Application US/09656002

Sequence 1, Application US/09656002

Sequence 1, Application US/09656002

GENERAL INFORMATION:

APPLICANT: Mack, David

APPLICANT: Milson, Keith

TITLE OF INVENTION: OVEREDATE CANCER MODULATORS

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AI

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER MODULATORS

FILE REFERENCE: A-69108/DJS/JJD/ANS

CURRENT APPLICATION NUMBER: US/09/656,002

CURRENT FILING DATE: 2000-03-16

PRIOR APPLICATION NUMBER: US/09/656,002

CURRENT FILING DATE: 2000-03-15

PRIOR FILING DATE: 2000-03-15

PRIOR FILING DATE: 2000-03-15

NUMBER OF SEQ ID NOS: 3

SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: PATENTIN VARIENT: A SOFTWARE: A S
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                                            AspLysLeuGlySerPheProSerLeuAlaValAlaLysIleIleIleIleGluPheAsm
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CRGANISM: Homo sapiens
US-09-656-002-1
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                                                                                                                                                                                                                            LIBRARY: COLNNOT13
CLONE: 1337018
SEQUENCE DESCRIPTION: SEQ ID NO: 18:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
TELEPAX: 650-845-4166
INFORMATION FOR SEQ ID NO: 18: SEQUENCE CHARACTERRISTICS:
LENGTH: 2038 base pairs
                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
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APPLICANT: Tavtigian, Sean V.
APPLICANT: Tavtigian, Sean V.
APPLICANT: Tavtigian, Sean V.
APPLICANT: Tavtigian, Sean V.
TITLE GNT: Myriad Genetics, Inc.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor
FILE REFERENCE: 2318-202
CURRENT APPLICATION NUMBER: US/09/342,749
CURRENT FILING DATE: 1999-06-29
EARLIER APPLICATION NUMBER: US 60/091,044
EARLIER PILING DATE: 1998-06-29
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PATENTI VEY: 2.0
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APPLICANT: Tavtigian, Sean V.
APPLICANT: Tang, David H.-F.
APPLICANT: Treng, David H.-F.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor
FILE REFERENCE: 2318-202
CURRENT APPLICATION NUMBER: US/09/691,840
CURRENT FILING DATE: 2000-10-18
PRIOR APPLICATION NUMBER: US/09/342,749
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                                                                                                                                     OTHER INFORMATION: This base can be C or T with C being the more OTHER INFORMATION: common allele. The codon is unaffected with both OTHER INFORMATION: alleles encoding Gly.
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                          ProSerThrProGlyValTyrThrLysValSerAlaTyrLeuAsnTrpIleTyrAsnVal 430
                                                                                                                                                                                                                                                                                           392 GlnSerAspGln---TrpHisValValGlyIleValSerTrpGlyTyrGlyCysGlyGly 410
                                                                      332 GlnAsnGlyGlyLysMetSerAspIleLeuLeuGlnAlaSerValGlnValIleAspSer 351
                                                                                                                                                                                                                     GlylleProGluGlyGlyValAspThrCysGlnGlyAspSerGlyGlyProLeuMetTyr 391
PheaspGluGluLeuThrProAlaThrProLeuTrpIleIleGlyTrpGlyPheThrLys 331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: conflict
LOCATION: (1471)
OTHER INFORMATION: Listed as A in Genbank Accession No. 6166194 U75329
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                                                                                                                                         352 ThrangCysasnalaaspaspalaTyrGinGlyGluValThrGluLysMetMetCysAla
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wong, Alexander K.C.
APPLICANT: Tavitigian, Sean V.
APPLICANT: Tavitigian, Sean V.
APPLICANT: Teng, DAvid H.-F.
APPLICANT: Wriad Genetics, Inc.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor
FILE REFERENCE: 2318-202
CURRENT PELLICATION NUMBER: US/09/342,749
CURRENT FILING DATE: 1999-06-29
EARLIER APPLICATION NUMBER: US 60/091,044
EARLIER PILING DATE: 1998-06-29
NUMBER OF SEQ ID NOS: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/09342749
Patent No. 6166194
GENERAL INFORMATION:
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SEQ ID NO 1
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ORGANISM: Homo sapiens
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NAME/KEY: conflict
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LOCATION: (1)
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LOCATION:
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LOCATION:
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                                                                                                                                                                                                                                                                                                                                                   LOCATION: (1471)
O'THER INFORMATION: Listed as A in GenBank Accession No. 6444419 U75329.
O'THER INFORMATION: Listed as A in GenBank Accession No. 6444419 U75329.
NAME/KEY: allele
LOCATION: (478)
O'THER INFORMATION: This base can be G or A with G being the more
O'THER INFORMATION: Common allele. The codon will change from Val to
O'THER INFORMATION: Met.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            115 AspSerAlaThrGlyAsnTrpPheSerAlaCysPheAspAsnPheThrGluAlaLeuAla
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                                                                                                                          6444419 U75329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: allele
COGATION: (777)
OTHER INFORMATION: This base can be C or T with C being the more
OTHER INFORMATION: common allele. The codon is unaffected with k
OTHER INFORMATION: alleles encoding Gly.
NAME/KEY: allele
LOCATION: (768)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This base can be C or T with C being the more common allele. This is a silent polymorphism.
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                                       Listed as T in GenBank Accession NO.
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Conservative:
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                                                                                        LOCATION: (985)
OTHER INFORMATION: L.
NAME/KEY: conflict
COCATION: (1347)
COTHER INFORMATION: L.
NAME/KEY: conflict
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OTHER INFORMATION:
NAME/KEY: conflict
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OTHER INFORMATION:
OTHER INFORMATION:
   LOCATION: (724)
OTHER INFORMATION:
                                                                conflict
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   CTGCACGTCCACAACGTCCACGTGTGCGGAGGCTCCATCATCACCCCCCGAGTGGATCGTG 876
                                                                                                                                                                              -- ProSerLeuAlaValAlaLysIle 274
                                                                                                                                                                                                                                                                                                                                                                                                                   MetLysLeuGlnPheProLeuThrPheSerGlyThrValArgProlleCysLeuProPhe 311
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                                                                                                                                                                                                                  937 GGGATTTTGAGA---CAATCTTTCATGTTCTATGGAGCCGGATACCAAGTAGAAAAGTG
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                                                                ThralaalaHisCysPheArgLysHis---ThraspValPheAsnTrpLysValArgAla
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Patent No. 644413

GENERAL INFORMATION:
APPLICANT: Wong, Alexander K.C.
APPLICANT: Tavtigian, Sean V.
APPLICANT: Tartigian, Sean V.
APPLICANT: Tartigian, Sean V.
APPLICANT: Myriad Genetics, Inc.
ITLLE OF INVENTION: TWRRS2 is a Tumor Suppressor FILE REFERENCE: 2318-202
CURRENT FALLIAG DATE: 2000-10-18
PRIOR APPLICATION NUMBER: US/09/691,840
CURRENT FILING DATE: 1999-06-29
PRIOR PLING DATE: 1999-06-29

PRIOR PLING DATE: 1999-06-29

PRIOR FILING DATE: 1998-06-29
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1462 ATGAGGGCAGAC 1473
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SEQ ID NO 1
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ORGANISM: Homo sapiens
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277 IleGluPheAsnProMetTyr---ProLys-----AspAsnAspIleAlaLeuMetLys 293
    ORGANISM: Homo sapiens
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CTGCACGTCCAGAACGTCCACGTGTGCGGAGGCTCCATCATCACCCCCGAGTGGATCGTG
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TCGAAGAATATCTGGTGGCTGATAGGGGATACAAGCTGGGGGTTCTGGCTGTGCCAAA
TCAAAAGCAGTGGTTTCTTTACGCTGTATAGCCTGCGGGGGTCAACTTGAACTCAAGCCGC
                                          ThrProArgValValGlyGlyGluGluAlaSerValAspSerTrpProTrpGlnValSer
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                                                                       ACAGOCOCOCACTGOGTAGAACCTCTTAACAATCCATGGCATTGGACGGCATTTGCG
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APPLICANT: O'Brien, Timothy J.
APPLICANT: O'Brien, Timothy J.
TITLE O'Brien, Transmembrane Serine Protease Overexpressed
TITLE OF INVENTION: In Ovarian Carcinoma and Uses Thereof
FILE REFERENCE: D6192CTP
CURRENT APPLICATION NUMBER: US/09/518,046
CURRENT FILING DATE: 2000-03-02
EARLIER FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 153
SEQ ID NO 1
LENGTH: 2413
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Matches:
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Mismatches:
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; FEATURE:
; NAMBKEY: CDS
: OTHER INFORMATION: entire CDNA sequence of
US-09-518-046-1
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52.14%
35.44%
27.78%
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us-09-607-745-2.rni

AlaSerileileileValValValLeuileLysValileLeuAspLysTyrATATTAGCACTGGCATTGGTCTGGGCATCGACTGCTCAGGGAGTACATATTAGCACTGGCATTGGTCTGGGCATCGCTCGGCTGGCT	468		222 TyrAspLysGlnHisValCysGlyGlySerIleLeuAspProHisTrpValLeuThrAla 241 852 TYCCAGGGCTACCACCTGTGGGGGGCTCTGTCATCACGCCCTGTGGATCATCACTGCT 911 242 AlaHis	1031 CCAGTGGACCATGTTATGGCAGATACAGGCTTGTACCTCCCCAAGTCATGGACCATCCAG 1090 258 AlaGlySerAspLysLeuGlySerPheProSerLeuAlaValAlaLysIleIle 275 1091 GTGGGTTACCTGTTGGACAATCCAGCCCATCCCACTTGGTGGACAAGATTGTC 1150 276 IlelleGluPheAsnProMetTyrProLysAspAsnAspIleAlaLeuMet 292 1151	1151TACCACAGCAAGTÁCAAGCCTÁAÁGAGGCTGGGCÁÁTGÁCCCTTÁTG 1201 293 LysLeuGlnPheProLeuThrPheSerGlyThvalArgProIleCysLeuProPhePhe 312
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	CyshanalahspaspalaTyrGlnGluValThrGluLysMetMetCyshlaGlylle	GGTGTACACCGTGTCACCTCCTTCCTGGACTGGATCCACGAGCAGATG u 435 h 1499 ion US/09518046	GENERAL INFORMATION: APPLICANT: O'Brien, Timothy J. APPLICANT: O'Brien, Timothy J. ITILE OF INVENTION: Transmembrane Serine Protease Overexpressed ITILE OF INVENTION: In Ovarian Carcinoma and Uses Thereof ITILE OF INVENTION: In Ovarian Carcinoma and Uses Thereof CURRENT FILE REPRENCE: D6192CIP CURRENT APPLICATION NUMBER: US/09/518,046 GURENT TAILING DATE: 2000-03-02 FARLIER PELLING DATE: 1999-03-03 NUMBER OF SEQ ID NOS: 153 SEQ ID NO 3 LENGTH: 2544 TYPE: DNA TYPE: DNA TYPE: DNA	NA sequence of TADG-12 Length: Matches:	nt Similarity: 48.14* Local Similarity: 33.06* Match: 3.06* 3.06* -607-745-2 (1-435) x US-09-518-04(22 IleproMetGluThrPheArgLys' ::: ::: 273 CTGCCATTTGAAGTTTTTCCCAA

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    -GICCGGGIGGGIGGICAGAAIGCCGIGCICCAGGIGIIC 506
                                115 AspSerAlaThrGlyAsnTrpPheSerAlaCysPheAspAsnPheThrGluAlaLeuAla 134
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               GATGGAGGTGACGCCTCCCCTGTCCTGAACCACGCGGCCGTCCCTTTGATTTCCAACAAG 1378
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Sequence 1. Application US/09261416A
Patent NO. 6291663
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Underwood, Lowell J.
TITLE OF INVENTION: Overexpressed in Ovarian Carcinoma
FILE REFERENCE: D6192
CURRENT APPLICATION NUMBER: US/09/261,416A
CURRENT APPLICATION NUMBER: US/09/261,416A
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 14
SEQ ID NO
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ORGANISM: Homo sapiens
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OTHER INFORMATION: CDS
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Best Local Similarity:
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; Patent No. 6518028
; GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
; TITLE OF INVENTION: Compositions and Methods
; TITLE OF INVENTION: O'WATAN CANCER
; TITLE OF INVENTION: O'WATAN CANCER
; TITLE OF INVENTION: O'WATAN CANCER
; CURRENT APPLICATION NUMBER: US/09/861,966
; CURRENT FILING DATE: 2001-05-21
; PRIOR APPLICATION NUMBER: 09/510,738
; PRIOR FILING DATE: 2000-02-22
; NUMBER OF SEQ ID NOS: 188
; SEQ ID NO 188
; LENGTH: 1783
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                                  Sequence 188, Application US/09510738A
Parent No. 6268165
GRNERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
TITLE OF INVENTION: Compositions and Methods for the Early Diagnosis of TITLE OF INVENTION: Ovarian Cancer
TITLE OF INVENTION: Ovarian Cancer
TITLE OF INVENTION: US/09/510,738A
CURRENT APPLICATION NUMBER: 105/09/510,738A
CURRENT FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: 09/039,211
PRIOR FILING DATE: 03-14-1998
NUMBER OF SEQ ID NOS: 188
SEQ ID NOS: 188
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ORGANISM: Homo sapiens
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Best Local Similarity:
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                                 32 ATCGTGGGAGGCCGGGACACCAGCTTGGGCCGGTGGCCGTGGCAAGTCAGCCTTCGCTAT
                                                                                                                                                                                                                           HisCysPheArgLysHisThrAspValPheAsn---TrpLysValArgAlaGlySerAsp
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                                                                                                               AsplysGlnHisValCysGlySerIleLeuAspProHisTrpValLeuThrAlaAla
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ORGANISM: Homo sapiens
FORTUME:
OTHER INFORMATION: full length cDNA of hepsin
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1389 GCCCTGGCCCAGAAGCCAGGCGTCTACACCAAAGTCAGTGACTTCCGGGAAGTGGATCTTC 1448
                                                                                                           149 TATGGCCAACAGGCGGGGTACTCCAGGAGGCTCGAGTCCCCCATAATCAGCAATGATGTC 1208
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1032 CTCTCCAGTCCCCTGCCCCTCACAGAATACAFCCAGCCTGTGTGCCTCCCAGCTGCCGGC 1091
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                            314 GluGluLeuThrProAlaThrProLeuTrpIleIleGlyTrpGlyPheThrLysGlnAsn 333
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Patent No. 6423543
GENERAL INFORMATION:
APPLICANT: Patrick Allen Marcotte
APPLICANT: Patrick Allen Marcotte
TITLE OF INVENTION: ANTISENSE MODULATION OF HEPSIN EXPRESSION
FILE REFERENCE: RTS-0090
CURRENT APPLICATION UNMER: US/09/742,703
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 49
SEQ ID NO 3
LENGTH: 2363
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ORGANISM: Homo sapiens
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Best Local Similarity:
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COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTAARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/000,846
FILING DATE: 30-DEC-1997
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/866,058
FILING DATE: 30-MY-1997
ATTORNEY/AGENT INFORMATION:
NAME: LEBOVITZ, RICHARD M.
REGISTRATION NUMBER: BERLX 65P1
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches:
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TELEPAN: 703-243-6410
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1605 base pairs
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STRANDEDNESS: single
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MOLECULE TYPE: CDN
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ANTI-SENSE: NO
ORIGINAL SOURCE:
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          ---GATTGCCCCAGAGGCCGTTTCTTGGCCGCCATCTGCCAAGACTGTGGCCGCAGGAAG 525
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Patent No. 5981830
GENERAL INFORMATION
APPLICANT: WU, QINGYU
APPLICANT: SADLER, JASPER
TITLE OF INVENTION: KNOCKOUT MICE AND THEIR PROGENY WITH
TITLE OF INVENTION: DISRUPTED SERINE PROTEASE GENES
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: MILLEN, WHITE, ZELANO & BRANIGAN, P.C.
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CITY: ARLINGTON
STRATE: VA
COUNTRY: US
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COMPUTER READABLE FORM:
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US-09-000-846-1
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Best Local Similarity:
SEQ ID NO 80
LENGTH: 1460
TYPE: DNA
ORGANISM: Homo s
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uLysThrProArgValvalGlyGlyGluGluAlaSerValAspSerTrpProTrpGlnya
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APPLICANT: Reed, Steven G.
APPLICANT: Reed, Steven G.
APPLICANT: Reed, Steven G.
APPLICANT: Mohamath, Roadoh
APPLICANT: Mohamath, Roadoh
APPLICANT: Mohamath, Roadoh
APPLICANT: Mohamath, Roadoh
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 21012.4.4.28
CURRENT APPLICATION NUMBER: US 09/285,323
FARLIER APPLICATION NUMBER: US 09/285,323
FARLIER PILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                          296 PheProLeuThrPheSerGlyThrValArgProIleCysLeuProPheAspGluGlu 315
276 IlelleGluPheAsnProMetTyrProLysAspAsnAspIleAlaLeuMetLysLeuGln 295
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Search completed: December 2, 2003, 02:54:14 Job time : 107.769 secs

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Sequence Sequence Sequence

10-174-591-330

Sequence Sequence Sequence

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Sequence Sequence Sequence

Sequence

10-174-569-330

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Sequence 6, Application US/10180719;
Publication No. US20030166246A1
GENERAL INFORMATION:
APPLICANT: BIllman, Jennifer L.
Yue, Henry
Guegler, Karl J.
Corley, Neil C.
Tang, Tom Y.
Shah, Purvi
TITLE OF INVENTION: HUMAN PROTEASE MOLECULES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Mohan-Peterson, Sheela
REGISTRATION NUMBER: 41,201
REFERENCE/DOCKET NUMBER: PF-0458 US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/180,719
FILING DATE: 25-Um-2002
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/008,271
FILING DATE: 16-Jan-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 650-855-055
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                                      US-10-180-719-6
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equence 8, Appli
                                                                                                                  ; Search time 324.904 Seconds
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246.943 Million cell updates/sec
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(gun2_6/ptodata/2/pubpaa/PUSO7_NEW_PUB.pep:*

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enCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-946-275
US-10-063-734-275
US-10-063-735-112
US-11-109-672-330
US-10-109-672-330
US-10-104-457-330
US-10-194-457-330
US-10-194-457-330
US-10-196-741-330
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US-09-851-588-8
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| APPLICANT: Edgar O. Ong |
| APPLICANT: Edgar O. Ong |
| APPLICANT: Edgar O. Ong |
| APPLICANT: Edgar O. Ong |
| APPLICANT: Jiunn-Chenr Yeh |
| APPLICANT: Jiunn-Chenr Yeh |
| APPLICANT: Jiunn-Chenr Yeh |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING |
| PRIOR APPLICATION NUMBER: 60/213,124 |
| PRIOR PELING DATE: 2000-06-22 |
| PRIOR PELING DATE: 2000-06-22 |
| PRIOR PELING DATE: 2000-06-21 |
| PRIOR APPLICATION NUMBER: 60/183,542 |
| PRIOR APPLICATION NUMBER: 60/183,542 |
| PRIOR APPLICATION NUMBER: 09/657,968 |
| PRIOR APPLICATION NUMBER: 00/657,968 |
| PRIOR APPLICATION NUMBER: 109/657,968                                                             99.8%; Score 2337; DB 9; I
100.0%; Pred. No. 6.7e-226;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99.08;
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                                                         Query Match
Best Local Similarity 100.
Matches 434; Conservative
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ORGANISM: Homo Sapien
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Sequence 8, Application US/09851588
Fatent No. US2020042067A1
GENERAL INFORMATION:
REPLICANT: Mack, David
APPLICANT: Mack, David
APPLICANT: Wilson, Keith E.
TITLE OF INVENTION: OF SCRENING FOR COLORECTAL CANCER, COME
TITLE OF INVENTION: OF SCRENING FOR COLORECTAL CANCER, COME
TITLE OF INVENTION: OF SCRENING FOR COLORECTAL CANCER, COME
TITLE OF INVENTION: OF SCRENING FOR COLORECTAL CANCER MODULATORS
FILE REPERENCE: A-68829-1/DJB/JJD/AMS
CURRENT APPLICATION NUMBER: US 09/642,552
FRIOR TILING DATE: 2000-09-17
FRIOR APPLICATION NUMBER: US 09/656,002
FRIOR FILING DATE: 2000-09-06
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.1
SEQ ID NO 8
LENGTH: 437

WANTER DEAL
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Pred. No. 5.3e-226;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                   9
                                                                                                                                                                                                                                                                        ; SEQUENCE DESCRIPTION: SEQ ID NO: US-10-180-719-6
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERIZIES: LENGTH: 435 amino acids
                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: COLNNOTI3
                                                                                                                                                                                                                                                                                                                                                              99.8%;
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ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-851-588-8
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299 GTVRPICLPFFDEBLTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAYQ 358
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                                                                                                                                                                                                                                                                                                                                                                                                                             Score 2297.5; DB 11; Length 432;
Pred. No. 6.1e-222;
0; Mismatches 0; Indels 5;
                  PRIOR FILING DATE: 2000-03-30
PRIOR APPLICATION NUMBER: PCT/USO0/23328
PRIOR APPLICATION NUMBER: PCT/USO0/32678
PRIOR FILING DATE: 2000-12-01
PRIOR FILING DATE: 2000-12-01
PRIOR PILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: PCT/USO1/06520
PRIOR APPLICATION NUMBER: PCT/USO1/06666
PRIOR APPLICATION NUMBER: PCT/USO1/06666
NUMBER OF SEQ ID NOS: 10
     APPLICATION NUMBER: PCT/US00/08439
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Grimaldi, Christopher J.
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Raton, Dan L.
Ferrara, Napoleone
Fong, Sherman
Gao, Wei-Qiang
Goddard, Audrey
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98.8%;
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Hillan, Kenneth J.
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Paoni, Nicholas F.
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Botstein, David
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 98.8
Matches 429; Conservative
                                                                                                                                                                                                                                                                                         ; LENGTH: 432
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-888-257A-7
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                                                                                                                           DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL 63
                                                                                                                                                                                                                                                                                 122 FSACFDNFTEALAETACROMGYSSKPTFRAVEIGPDODLDVVEITENSOELRMRNSSGPC
                                                                                DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL
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                              Gaps
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APPLICANT: Hillar, Kenneth J.
APPLICANT: Smith, Victoria
APPLICANT: Work, William I.
APPLICANT: Work, William I.
APPLICANT: Wu, Thomas D.
APPLICANT: Chang, Zemin I.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND TITLE OF INVENTION: TREATMENT OF TUMOR
FILE REPERENCE: PSOORI
CURRENT APPLICATION NUMBER: US/09/888,257A
CURRENT PILLING DATE: 2002-04-30
                              0;
                              Indels
     Pred. No. 4.3e-224;
                              0; Mismatches
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PRIOR PILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: US 60/089,653
PRIOR PILING DATE: 1998-06-17
PRIOR FILING DATE: 1998-06-17
PRIOR PLING DATE: 1998-06-10
PRIOR PLING DATE: 1998-06-10
PRIOR APPLICATION NUMBER: US 60/13,678
PRIOR APPLICATION NUMBER: US 60/235,451
PRIOR APPLICATION NUMBER: PCT/US99/12252
PRIOR PRILING DATE: 1998-10-08
PRIOR PILING DATE: 1998-10-08
PRIOR PILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: PCT/US99/20111
PRIOR APPLICATION NUMBER: PCT/US09/20111
PRIOR PILING DATE: 1999-09-01
PRIOR PILING DATE: 1999-09-01
PRIOR PILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
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     99.5%;
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APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
                              432; Conservative
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-888-257A-7
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R FILING DATE: 1998-10-01
R APPLICATION NUMBER: 60/102965
R APPLICATION NUMBER: 60/103288
R FILING DATE: 1998-10-05
R FILING DATE: 1998-10-06
R FILING DATE: 1998-10-07
R FILING DATE: 1998-10-07
R FILING DATE: 1998-10-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R FILING DATE: 1998-09-30
R APPLICATION NUMBER: 60/102571
R FILING DATE: 1998-09-30
R APPLICATION NUMBER: 60/102684
R FILING DATE: 1998-10-01
R APPLICATION NUMBER: 60/102687
                       PPLICATION NUMBER: 60/100711
LLING DATE: 1998-09-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 1998-09-30
APPLICATION NUMBER: 60/102487
FILING DATE: 1998-09-30
APPLICATION NUMBER: 60/102570
                                 PRIOR 
                                                                                     PRIOR APPLICATION NUMBER: 60/100684
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: 60/100710
                Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
Watanabe, Colin K.
Williams, P. Mickey
Wood, William I.
     Roy, Margaret Ann
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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302 GTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAYQ 361
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Gao, Wel-Giang
APPLICANT: Godowski, Audrey
APPLICANT: Godowski, Audrey
APPLICANT: Godowski, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Hillan, Kenneth J.
APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Pan, Micholas F.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILLE OF INVENTION: 2014 Secreted and Transmembrane Polypeptides and Nucleic
FILLE PREFERENCE: P283 OPLICA: 12-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             242 AHCFRKHTDVFNWKVRAGSDKLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTFS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   359 GEVIEKMMCAGIPEGGVDTCQGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prior Application removed - See File Wrapper or Palm NUMBER OF SEQ ID NOS: 477 SEQ ID NO 275
                                                                                                     Sequence 275, Application US/10015387A Publication No. US20030135034A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                    Ferrara, Napoleone
Fong, Sherman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    422 AYLNWIYNVWKAEL 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             419 AYLINMIYNVWKAEL 432
                                                                                                                                                                                                    APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eaton, Dan 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 98.8
Matches 429; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-015-387A-275
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US-10-063-735-112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL
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Pred. No. 6.1e-222;
0; Mismatches 0;
PRIOR PILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: 60/10335
PRIOR APPLICATION NUMBER: 60/10336
PRIOR PILING DATE: 1998-10-07
PRIOR PELING DATE: 1998-10-07
PRIOR PILING DATE: 1998-10-07
PRIOR PILING DATE: 1998-10-06
PRIOR PILING DATE: 1998-10-06
PRIOR APPLICATION NUMBER: 60/103633
PRIOR PILING DATE: 1998-10-08
PRIOR PILING DATE: 1998-10-07
PRIOR PILING DATE: 1998-10-20
PRIOR APPLICATION NUMBER: 60/105104
PRIOR APPLICATION NUMBER: 60/10509
PRIOR PILING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: 60/105169
PRIOR PILING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: 60/105694
PRIOR PILING DATE: 1998-10-22
PRIOR APPLICATION NUMBER: 60/105694
PRIOR PILING DATE: 1998-10-22
PRIOR PILING DATE: 1998-10-22
PRIOR PILING DATE: 1998-10-22
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PRIOR APPLICATION NUMBER: 60/105807
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98.8%;
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Best Local Similarity
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APPLICANT: Pan, James
APPLICANT: Pan, James
APPLICANT: Panni, Nicholas F.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
TITLE OF INVENTION: Acids Encoding the Same
TITLE OF INVENTION: Acids Encoding the Same
CURRENT APPLICATION NUMBER: US/10/006,130A
CURRENT APPLICATION NUMBER: US/10/006,130A
PITOR APPLICATION TEMOVED - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 477
SEQ ID NO 275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 299 GTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAYQ
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Pred. No. 6.1e-222;
0; Mismatches 0; Indels
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GENERAL INFORMATION: APPLICANT: BAKET, KEVIN P. APPLICANT: Chen, Jian
                                                    Godowski, Paul J.
Grimaldi, Christopher J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98.1%;
98.8%;
                                                                                                  Gurney, Austin L.
Hillan, Kenneth J
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Wood, William I.
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                          Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 98.8
Matches 429; Conservative
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Smith, Victoria
       Wei-Qiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Desnoyers, Luc
Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-006-130A-275
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US-10-199-672-330
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APPLICANT: I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          63
                                                                                                                                  APPLICANT: GGALTASEL, MALY B.
APPLICANT: GGALACEA, MALEY
APPLICANT: GGAGARA, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Grimalos, Colin E.
APPLICANT: Watanabe, Colin E.
APPLICANT: Watanabe, Colin E.
APPLICANT: Wood, William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3230RIC1
CURRENT APPLICANTION NUMBER: US/10/063,735
CURRENT FILING DATE: 2002-05-08
PRIOR APPLICANTION TEMOVED - See Palm or File Wrapper
NUMBER OF SEQ ID NOS: 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 2297.5; DB 12; Length 432; Pred. No. 6.1e-222;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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Sequence 112, Application US/10063735
Publication No. US20030138882A1
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Eston, Dan I.
APPLICANT: Ferrara, Napoleone
APPLICANT: Forg, Sherman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 98.8%;
Matches 429; Conservative (
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                                                            APPLICANT: Eaton, Dan L.
APPLICANT: Filvaroff, Ellen
APPLICANT: Gerritsen, Mary E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-063-735-112
                        Publication No. US20
GENERAL INFORMATION:
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LENGTH: 432
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63

Gaps

181

241

301 298 361 358 421

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APPLICANT: Paoni, Nicholas F. IIILE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILE REFERENCE: P2830P1011
CURRENT APPLICATION NUMBER: US/10/006,172A
CURRENT FILING DATE: 2002-03-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/09116
PRIOR PILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-01
PRIOR PLING DATE: 1998-09-01
PRIOR PLING DATE: 1998-09-01
PRIOR PELING DATE: 1998-09-01
PRIOR PELING DATE: 1998-09-02
PRIOR PILING DATE: 1998-09-02
PRIOR PELING DATE: 1998-09-09
PRIOR PELING DATE: 1998-09-10
                                                                                                WT: Baker, Kevin P.
WT: Betstein, David
WT: Desnoyers, Luc
MT: Eaton, Dan 1.
MT: Ferrara, Napoleone
MT: Fong, Sherman
MT: Gao, Wei-Qiang
MT: Godaxd, Audrey
MT: Godowski, Paul J.
MT: Grimaldi, Christopher J.
Sequence 275, Application US/10006172A Publication No. US20030153000A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 1998-09-16
APPLICATION NUMBER: 60/100627
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APPLICATION NUMBER: 60/100661
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Hillan, Kenneth J
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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                     TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REPERENCE: P3430R1C1
CURRENT APPLICATION NUMBER: US/10/199,672
CURRENT FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: US/10/052,586
PRIOR PILING DATE: 1997-09-18
PRIOR PILING DATE: 1997-09-18
PRIOR PILING DATE: 1997-09-18
PRIOR PILING DATE: 1997-09-18
PRIOR PELING DATE: 1997-09-18
PRIOR PELING DATE: 1997-10-17
PRIOR PELING DATE: 1997-10-24
PRIOR PELING DATE: 1997-10-24
PRIOR PILING DATE: 1997-10-28
PRIOR PELING DATE: 1997-10-28
PRIOR PILING DATE: 1
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Pred. No. 6.1e-222;
0; Mismatches 0; Indels 5;
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98.8%;
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Best Local Similarity 98.8
Matches 429; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
CRGANISM: Homo Sapien
US-10-199-672-330
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US-10-006-172A-275
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PRIOR FILING DATE: 1998-09-16
PRIOR APPLICATION NUMBER: 60/100662
PRIOR PLILING DATE: 1998-09-16
PRIOR PLILING DATE: 1998-09-16
PRIOR FILING DATE: 1998-09-16
PRIOR FILING DATE: 1998-09-17
PRIOR PLILING DATE: 1998-09-18
PRIOR PLILING DATE: 1998-09-17
PRIOR PLILING DATE: 1998-09-18
PRIOR PLILING DATE: 1998-09-17
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PRIOR PLILING DATE: 1998-09-17
PRIOR PLILING DATE: 1998-09-18
PRIOR PLILING DATE: 1998-09-23
PRIOR PLILING DATE: 1998-09-24
PRIOR PLILING DATE: 1998-09-24
PRIOR PLILING DATE: 1998-09-24
PRIOR PLILING DATE: 1998-09-29
PRIOR PLILING DATE: 1998-09-3
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LSGSLVSLHCLACGKSLKTPRVVGGEBASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTA 241 64 CGQPLHFIPRKQLCDGBLDCPLGEDEBHCVKSFFBGPAVAVRLSKDRSTLQVLDSATGNW 123 122 FSACFDNFTEALAETACRQMGYSSKPTFRAVEIGPDQDLDVVEITENSQELRMRNSSGPC 181 124 FSACFDNFTEALAETACROMGYS-----RAVEIGPDQDLDVVEITENSQELRMRNSSGPC 178 4 DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL 2 DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL 62 CGQPLHFIPRKQLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNW 5. Query Match 98.1%; Score 2297.5; DB 12; Length 432; Best Local Similarity 98.8%; Pred. No. 6.1e-222; Matches 429; Conservative 0; Mismatches 0; Indels 5; PRIOR APPLICATION NUMBER: 60/102687
PRIOR FILING DATE: 1998-10-01
PRIOR FILING DATE: 1998-10-02
PRIOR FILING DATE: 1998-10-02
PRIOR FILING DATE: 1998-10-05
PRIOR PELING DATE: 1998-10-07
PRIOR PELING DATE: 1998-10-08
PRIOR PELING DATE: 1998-10-20
PRIOR PELING DATE: 1998-10-21
PRIOR PELING DATE: 1998-10-22
PRIOR PELING DATE: 1998-10-27
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US-10-194-457-330
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APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul J.
APPLICANT: Gurney, Austin L.
APPLICANT: Ban, James
APPLICANT: Matanabe, Colin K.
APPLICANT: Matanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Mod William I.
APPLICANT: NORE BNCODING THE SAME
ITILE OF INVENTION: SECRETED AND TRANSMENBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: SCRETED AND TRANSMENBRANE
TITLE APPLICATION NUMBER: US/10/187,749
CURRENT FILING DATE: 2002-07-01
PRIOR APPLICATION NUMBER: GO/05266
PRIOR PILING DATE: 1997-09-18
PRIOR APPLICATION NUMBER: GO/05266
PRIOR PILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: GO/063120
PRIOR PILING DATE: 1997-10-24
PRIOR APPLICATION NUMBER: GO/06346
PRIOR FILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: GO/06344
PRIOR PILING DATE: 1997-10-28
PRIOR PILING DATE: 1997-10-28
PRIOR PILING DATE: 1997-10-28
PRIOR APPLICATION NUMBER: GO/063541
PRIOR PILING DATE: 1997-10-28
PRIO
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LSGSLVSLHCLACGKSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTA 238
                                                          242 AHCFRKHTDVFNWKVRAGSDKLGSFPSLAVAKIIIIBFNPMYPKDNDIALMKLQFPLTFS 301
                                                                                                   AHCERKHTDVFNWKVRAGSDKLGSFPSLAVAKIIIEFNPMYPKDNDIALMKLQFPLTFS
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Pred. No. 6.1e-222;
0; Mismatches 0; Indels 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 330, Application US/10187749 Publication No. US20030153036A1 GENERAL INFORMATION:
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Best Local Similarity 98.8%;
Matches 429; Conservative
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
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; ORGANISM: Homo Sapien
US-10-187-749-330
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APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Smith, Victoria
APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colin K.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACLDS ENCODING THE SAMB
TITLE OF INVENTION: ACLDS ENCODING THE SAMB
FILE REPERENCE: 1202-07-11
FILE REPERENCE: 2002-07-11
FRICK APPLICATION NUMBER: 10/05286
FRICK APPLICATION NUMBER: 60/05266
FRICK APPLICATION NUMBER: 60/05266
FRICK APPLICATION NUMBER: 60/062250
FRICK APPLICATION NUMBER: 60/062120
FRICK APPLICATION NUMBER: 60/063120
FRICK APPLICATION NUMBER: 60/063120
FRICK APPLICATION NUMBER: 60/063121
FRICK APPLICATION NUMBER: 60/063121
FRICK APPLICATION NUMBER: 60/063121
FRICK APPLICATION NUMBER: 60/063121
FRICK APPLICATION NUMBER: 60/063146
FRICK FILING DATE: 1997-10-21
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                                                                                                                       CGQPLHFIPRKQLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNW 121
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Sequence 330, Application US/10194457
Publication No. US20030153037A1
GENERAL INFORMATION:
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APPLICATION NUMBER: 60/063541
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APPLICATION NUMBER: 60/063544
FILING DATE: 1997-10-28
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Gurney, Austin L.
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APPLICANT: Smith, Victoria
APPLICANT: Matanabe, Colin K.
APPLICANT: Matanabe, Colin K.
APPLICANT: Mood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
TITLE OF INVENTION: ACIDS ENCODING THE SAME
CURRENT PELING DATE: 2012-06-27
Prior Application removed - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 612
                                                                                                                                                                                                                                  4 DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL
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                                                                                                                                          Length 432;
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 - See File Wrapper or
                                                                                                                                         DB 12;
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Pred. No. 6.1e-222;
0; Mismatches 0;
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Publication No. US20030157635A1
GENERAL INFORMATION:
Prior Application data removed
NUMBER OF SEQ ID NOS: 612
SEQ ID NO 330
                                                                                                                                       98.1%;
98.8%;
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Gurney, Austin L.
Pan, James
                                                                                                                                                                            Matches 429; Conservative
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Goddard, Audrey
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US-10-184-642-330
                                                                     TYPE: PRT
CORGANISM: Homo Sapien
US-10-194-457-330
                                                                                                                                       Query Match
Best Local Similarity
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US-10-184-642-330
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APPLICANT: Wood William I.
APPLICANT: Wood William I.
APPLICANT: Wood William I.
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REPREBNCE: 1943.046
CURRENT APPLICATION NUMBER: US/10/196,747
CURRENT FILING DATE: 2002-07-16
                                                                                                                                                                                                                                                                                     62 CGQPLHFIPRKQLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNW
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  432;
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  Length
Score 2297.5; DB 12; Lengt
Pred. No. 6.1e-222;
0; Mismatches 0; Indels
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NUMBER OF SEQ ID NOS: 612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 330, Application US/10196747
Publication No. US20030162250A1
GENERAL INFORMATION:
98.1%;
98.8%;
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Smith, Victoria
Watanabe, Colin K.
Wood, William I.
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Best Local Similarity 98.8
Matches 429; Conservative
                 al Similarity 98.8
429; Conservative
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Goddard, Audrey
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; ORGANISM: Homo Sapien
US-10-196-747-330
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US-10-196-747-330
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Query Match
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   PRIOR APPLICATION NUMBER: 60/099598
PRIOR FILING DATE: 1998-09-09
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 477
SEQ ID NO 275
LENGTH: 432
                                                                                                                                                                                                                                                                                                                                                                                          DB 12; Length 432;
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Pred. No. 6.1e-222;
0; Mismatches 0; Indels
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ilarity 98.8%;
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 429; Conserv
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                                                                                 CGOPLHFIPRKOLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNW 121
                                                                                                                                                    CGQPLHFIPRKQLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNW 123
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4 DPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFL 63
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APPLICANT: Hillan, Kenneth J.
APPLICANT: Pan, James A.
APPLICANT: P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/098716
PRIOR FILING DATE: 1996-09-01
PRIOR PELING DATE: 1996-09-01
PRIOR PELING DATE: 1996-09-01
PRIOR PELING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: 60/098749
PRIOR PILING DATE: 1998-09-01
PRIOR FILING DATE: 1998-09-02
PRIOR PELING DATE: 1998-09-02
PRIOR PELING DATE: 1998-09-02
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Fong, Sherman
Gao, Wel-Qiang
Goddard, Audrey
Godowski, Paul J.
Grimaldi, Christopher J.
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APPLICATION NUMBER: 60/099536
FILING DATE: 1998-09-09
APPLICATION NUMBER: 60/099596
FILING DATE: 1998-09-09
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APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
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CLONE: 1337018
SEQUENCE DESCRIPTION: SEQ ID NO:
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STRANDEDNESS: single
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US-09-008-271A-6
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                                                                                   475
472.5
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                                                                  2003, 00:14:04 ; Search time 31.7125 Seconds (without alignments) 580.377 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Al
Sequence 3, Al
Sequence 67, 2
Sequence 26, 2
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Sequence 2, 7
Sequence 2, 7
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1 MDPDSDQPLNSLDVKPLRKP......VYTKVSAYLNWIYNVWKAEL 435
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Sequence 3,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/pcTUS_COMB.pep:*
 GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-65-002-2
US-09-61-840-2
US-09-61-840-2
US-09-51-840-2
US-09-261-416-2
US-09-261-416-2
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US-09-370-838-62
US-09-370-838-62
US-09-00-151-1
US-08-807-151-1
US-08-807-151-1
US-08-807-151-1
US-08-907-337-3
US-09-027-337-3
US-09-64-483-67
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Maximum Match 100%
Listing first 45 summaries
                                                 OM protein - protein search, using sw model
                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                          Issued_Patents_AA:*
                                                                                                                                                                                                                                                        Minimum DB seq length: 0 Maximum DB seq length: 20000000000
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Match Length
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482.5
478
                                                                                                                      Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
CITY: Palo Alto
    US-09-344-675-4
US-09-386-629-7
US-09-644-600-10
US-09-734-675-2
US-09-734-675-2
US-08-944-483-68
US-09-624-337-2
US-09-644-600-2
US-09-387-375-9
US-08-391-761A-13
US-08-451-932-1
US-08-451-932-1
US-08-451-932-1
US-08-451-932-1
US-08-451-932-1
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US-08-451-932-1
US-08-451-932-1
US-08-605-5988-1
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                    US-08-429-743-1
                                                                                                                                                                                                                     ALIGNMENTS
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INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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ZIP: 94304
COMPUTER READABLE FORM:
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435
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                                                                                                    184 KSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWK
                                                                                                                                                                                                                                                         244 VRASSDKLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEB
                                                                                                                                                                                                                                                                                                                                           304 LIPAIPLWIIGWGFIKQNGGKWSDILLQASVQVIDSTRCNADDAYQGEVIEKMMCAGIPE
                                                                                                                                                                                                                                                                                                                                                                                          376 GGVDTCQGDSGGPLMYQSDQWHVVGIVSMGXGCGGPSTPGVYTKVSAYLNWIYNVWKABL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKHTDVFNWK
                                                                                                                                                                                                                                  VRAGSDKLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGTVRPICLPFFDEE
                                                                                                                                                                                                                                                                                                                                                                                                                                 364 GGVDTCQGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKVSAYLNWIYNVWKAEL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                292 MKLQFPLTFSGTVRPICLPFFDBELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            352 TRCNADDAYQGEVTEKMMCAGIPEGGVDTCQGDSGGPLMYQSDQ-WHVVGIVSWGYGCGG
                                                                             TACROMGYSSKPTFRAVEIGPDQDLDVVEITENSQELRMRNSSGPCLSGSLVSLHCLACG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Wong, Alexander K.C.
APPLICANT: Trans, David H.-F.
APPLICANT: Trans, David H.-F.
APPLICANT: Myriad Genetics, Inc.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor;
FILE REPERBACE: 2318-202
CURRENT APPLICATION NUMBER: US/09/342,749
CURRENT FILING DATE: 1999-06-29
BARLIER FILING DATE: 1998-06-29
NUMBER OF SEC ID NOS: 33
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28.9%; Score 676.5; DB 3;
llarity 39.1%; Pred. No. 1.2e-60;
Conservative 57; Mismatches 128;
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Patent No. 6166194
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US-09-342-749-2
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Best Local Similarity
Matches 150; Conserv
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Sequence 2. Application US/09656002

Patent No. 645568

GENERAL INFORMATION:

APPLICANT: Milson, Keith

APPLICANT: Milson, Keith

TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING COLORECTAL CANCER, COMPOSITIONS, AND

TITLE OF INVENTION: NO SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AND

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AND

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AND

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AND

TITLE OF INVENTION: OF SCREENING FOR COLORECTAL CANCER, COMPOSITIONS, AND

FILE REFERENCE: A-69108/DISJAJANAS

CURRENT FILING DATE: 2000-09-15

PRIOR FILING DATE: 2000-01-28

PRIOR PRILING DATE: 2000-01-28

PRIOR APPLICATION NUMBER: US 09/493,444

PRIOR FILING DATE: 2000-03-15

NUMBER OF SEQ ID NOS: 3

SOFTWARE: PATENTING DATE: 2000-03-15

NUMBER OF SEQ ID NOS: 3

SOFTWARE: PATENTING DATE: 2000-03-15
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                                                                                                                                                                                                          61 LCGQPLHFIPRKQLCDGBLDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGN 120
                                                                                                                                                                                                                                                   121 WFSACPDNFTEALAETACROMGYSSKPTFRAVEIGPDQDLDVVEITENSQELRMRNSSGP 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SGTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAY 360
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                                                                                                                          MDPDSDQPLNSLDVKPLRKPRIPMBTFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SGTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGRWSDILLQASVQVIDSTRCNADDAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              361 QGEVTEKAMCAGIPEGGVDTCQGDSGGPLMYQSDQWHVVGIVSWGYGCGGPSTPGVYTKV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PLRKPRIPMETFRKVGIPILIALLSLASIIIVVVLIKVILDKYYFLCGQPLHFIPRKQLC
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                                                                                              MDPDSDQPLNSLDVKPLRKPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYF
                                                                                                                                                                                                                                                                                        121 WFSACFDNFTEALAETACROMGYSSKPTFRAVEIGPDODLDVVEITENSQELRMRNSSGP
                                                                                                                                                                                                                                                                                                                              CLSGSLVSLHCLACGKSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLT
                                                                                                                                                                                                                                                                                                                                                                                                            AAHCFRKHTDVFNWKVRAGSDKLGSFPSLAVAKIIIIEFNPMYPKDNDIALMKLQFPLTF
                                                         Gaps
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                  Length 435
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Pred. No. 2.1e-216;
1; Mismatches 11;
                99.8%; Score 2338; DB 3;
99.8%; Pred. No. 6.2e-231;
iive 1; Mismatches 0;
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Best Local Similarity 97.1%;
Matches 408; Conservative
          Query Match
Best Local Similarity 99.8
Matches 434; Conservative
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US-09-656-002-2
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LENGTH: 423
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INVENTION: Transmembrane Serine Protease Overexpressed INVENTION: in Ovarian Carcinoma and Uses Thereof
                                                             US/09/518,046
     TITLE OF INVENTION: Transmembrane Serine
TITLE OF INVENTION: in Ovarian Carcinoma
FILE REFERENCE: D6192CIP
CURRENT APPLICATION NUMBER: US/09/518,046
CURRENT FILING DATE: 2000-03-02
EARLIER FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 153
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Patent No. 6482630
GENERAL INFORMATION:
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Best Local Similarity 35.75
Matches 158; Conservative
                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
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LENGIH: 45
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APPLICANT: Wong, Alexander K.C.
APPLICANT: Tartigian, Sean V.
APPLICANT: Trang, David H.-F.
APPLICANT: Trang, David H.-F.
APPLICANT: Myriad Genetics, Inc.
TITLE OF INVENTION: TMPRSS2 is a Tumor Suppressor
FILE REPERENCE: 2318-202
CURRENT APPLICATION NUMBER: US/09/691,840
CURRENT APPLICATION NUMBER: US/09/342,749
PRIOR APPLICATION NUMBER: US/09/342,749
PRIOR APPLICATION NUMBER: US 60/091,044
PRIOR PILING DATE: 1998-06-29
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO S.
ILENGTH: 492
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                                                            434
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                                                                                          468 AYRPGVYGNVMVFTDWİYRQMRAD 491
                                                          411 PSTPGVYTKVSAYLNWIYNVWKAE
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Patent No. 6294663
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
APPLICANT: Underwood, Lowell J.
                                                                                                                                                                                  Sequence 2, Application US/09691840 Patent No. 6444419
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; ORGANISM: Homo sapiens
US-09-691-840-2
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APPLICANT: Differencesco, Valentina
APPLICANT: Differencesco, Valentina
APPLICANT: Beasley, Ellen
TITLE OF INVENTION: ISOLATED HUMAN PROTEASE PROTEINS,
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL001194
CURRENT APPLICATION NUMBER: US/09/820,002
CURRENT FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         311 LAGPLTENEMIQPVCLPNSEENFPDGKVCWTSGWGAT-EDGGDASPVLNHAAVPLISNKI 369
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                                                                                                                                                                                                                                                                22 IPMETFRKVGIPIIIALLSLASIIIVVVLIKVILD---KY----YFLCGQPLHFIPRKQL
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                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                          63;
                                                                                                                                                  Length 454.;
FEATURE:
OTHER INFORMATION: complete amino acid sequence of TADG-12
CTHER INFORMATION: protein
US-09-518-046-2
                                                                                                                                                                                                       Indels
                                                                                                                                            28.0%; Score 655.5; DB 3; 35.7%; Pred. No. 1.5e-58; iive 73; Mismatches, 149;
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Page

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APPLICANT: DiFrancesco, Valentina
APPLICANT: DiFrancesco, Valentina
APPLICANT: Beasley, Ellen
TITLE OF INVENTION: ISOLATED HUMAN PROTEASE PROTEINS,
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CLOO1194
CURRENT APPLICATION NUMBER: US/09/820,002
CURRENT APPLICATION NUMBER: 2001-03-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PastSEQ for Windows Version 4.0
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15 RPKVAALT---AGTLLLLTAIGAASWAIVAVLLR-
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Patent No. 6482630
GENERAL INFORMATION:
APPLICANT: Gan, Weiniu
APPLICANT: Ye, Jane
CDGVSDCKDGEDEYRC-
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                                                                                               ETACROMGYSS---
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TYPE: PRT
ORGANISM: HUMAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EEMĞF----LRAL---THSELDVRTAGANGTSGFFCVDEGRLPHTQRLLEVISPSDCPR 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        322 CNGADFYGNQIXPKMFCAGYPEGGIDACOGDSGGFFVCEDSISRIPRWRLCGIVSWGTGC 381
                                                                                                                                                                                                                                                                                                                                                                                LDCPLGEDERHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNWFSACFDNFTEALAETAC 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----SQELRMRNSSGPCLS 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSLVSLHCLACG-KSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAA 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HCFRKHTDVFN-WKVRAGSDKLGSFPSLAVAKIIIIEFNPMYP------KDNDIALMK 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       294 LOFPLIFSGIVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTR 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     263 LSSPLPLTEYIQPVCLPAAGQALVDGKICTVTGWGNT-QYYGQQAGVLQBARVPIISNDV 321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CNADDAYQGEVTEKMMCAGIPEGGVDTCQGDSGGPLMYQ----SDQWHVVGIVSWGYGC 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-261-416-2
Sequence 2, Application US/09261416A
Patent No. 6291663
GENERAL INFORMATION:
APPLICANT: O'Brien, Timothy J.
TITLE OF INVENTION: TADG-12: A No. 6291663el Transmembrane Serine Protease
TITLE OF INVENTION: Overexpressed in Ovarian Carcinoma
FILE REPRENCE: D6192
CURRENT APPLICATION NUMBER: US/09/261,416A
CURRENT APPLICATION NUMBER: US/09/261,416A
SEQ ID NO.2
SEQ ID NO.2
                                                                                                                                                                                                                                                                                                                                                                                                                           CDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNWFSACFDNFTEALA 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74
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Patent No. 6291663
98-09-261-416-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ----YFLCGQPLHFIPRKQL
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                                                                                                                                                                                                                                                      19 KPRIPMETFRKVGIPIIIALLSLASIIIVVVLIKVILDKYYFLCGQPLHFIPRKQLCDGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                   DB 4; Length 417;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 455;
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                                                                                                                         Similarity 31.5%; Pred. No. 9.7e-52; On Conservative 60; Mismatches 161;
                                                                                                                                                                                                                                                                                                    22 IPMETFRKVGIPIIIALLSLASIIIVVVLIKVILD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          139 ROMGYSSKPTFRAVEIGPDODLDVVEITEN--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  382 ALAQKPGVYTKVSDFREWIFQAIK 405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           109 GGPSTPGVYTKVSAYLNWIYNVWK 432
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 153; Conserv
                                HUMAN
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                             ; ORGANISM: HI
US-09-820-002-4
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                                                                                                                             Query Match
Best Local
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                                                 190 HCLACG-KSLKTPRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCFRKH 248
                                                                                                                                                                                                                                                                                                                                                                     327 YGNOIKPKWFCAGYPEGGIDACOGDSGGPFVCEDSISGTSRWRLCGIVSWGTGCALARKP 386
       -- CPRGRFLTA 147
                                                                                                                                                                     268 LTBYIQPVCLPAAGQALVDGKVCTVTGWGNT-QFYGQQAMVLQBARVPIISNEVCNSPDF
                                                                                                                                           249 TDVFN-WKVRAGSDKLGSFPSLAVAKIIIIEFNPMYP------KDNDIALMKLQFPLT
                                                                                                                                                                                                                                 300 FSGTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDA
                                                                                                                                                                                                                                                                                                                        YQGEVTEKMMÇAĞIPEĞĞVDIÇQĞDŞĞĞELMYQ----SDQWHVVĞIVŞWGYĞÇĞĞPSIP
                                                                                            148 TCQDCGRRKLPVDRIVGGQDSSLGRWPWQVSLRYDGTHLCGGSLLSGDWVLTAAHCFPER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69 IPRKOLCDGELDCPLGEDEBHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNWFSACFDN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                42; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLONING OF ENTEROXINASE AND METHOD OF USE
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Pred. No. 1.6e-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                67; Mismatches 138; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYGTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/200,900A
FILING DATE: 23-FEE-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B: Genetics Institute, Inc. - Legal Affairs
87 CambridgePark Drive
       101 SELDVRTAGANGTSGFFCVDEGGLRLAQRLLDVISVCD-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Meinert, Maureen C.
REGISTRATION NUMBER: 31,544
REFISTRATION NUMBER: 615201-FWC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 876-1170 X8574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/08200900A
Patent No. 566556
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: CLONING OF ENTE
NUMBER OF SEQUENCES: 38
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPAN: (617) 876-1170 X8574
TELEFAX: (617) 876-5851
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  GVYTKVSAYLNWIYNVWK 432
                                                                                                                                                                                                                                                                                                                                                                                                                                                             387 GVÝTKVTDFREWIFKAIK 404
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TYPE: amino acid
STRANDEDNESS: single
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Best Local Similarity 35.0%
Matches 133; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                                     296 FCAGYPEGGIDACQGDSGGPFVCEDSISRIPRWRLCGIVSWGTGCALAQKPGVYTKVSDF 355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CVKSFPEGPAVAVRLSKDRSTLQVLDSATGNWFSACFDNFTEALAETACRQMGY----- 143
                                                                                                                                         369 MCAGIPEGGVDTCQGDSGGPLMYQ----SDQWHVVGIVSWGYGCGGPSTPGVYTKVSAY 423
177 FAGAVAQASPHGLQLGVQAVVYHGGYLPFRDPNSBENSNDIALVHLSSPLPLTEYIQPVC 236
                                                 309 LPFFDEELTPATPLWIIGWGFTKONGGKMSDILLQASVOVIDSTRCNADDAYQGEVTEKM 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                48
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                                                                                              237 LPAAGQALVDGKICTVTGWGNT-QYYGQQAGVLQEARVPIISNDVCNGADFYGNQIKPKM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: SADLER, JASPER
TITLE OF INVENTION: KNOCKOUT MICE AND THEIR PROGENY WITH
TITLE OF INVENTION: DISRUPTED SERINE PROTEASE GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      62; Mismatches 151; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               : MILLEN, WHITE, ZELANO & BRANIGAN, P.C. 2200 CLARENDON BLVD. SUITE 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 574;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS SOFTMARE: PS ASTENTIN Release #1.0, Ver CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/000,846 FILING DATE: 30-DEC-1997 CLASSIFICATION: 800 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TRICKEL LEBOUITZ, RICHARD M.
REGISTRATION NUMBER: 37,067
REPERENCE/DOCKET NUBER: BERLK 65P1
TELECOMMUNICATION INFORMATION:
TRIEPHONE: 703-243-6333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/866,058
FILING DATE: 30-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 VGTLLFLTGIGAASWAİVTILLQ-
                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2, Application US/09000846
Patent No. 5981830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: WU, QINGYU APPLICANT: SADLER, JASPER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                              424 LNWIYNVWK 432
                                                                                                                                                                                                                                                                           356 REWIFQAIK 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC
OPERATING SYSTEM:
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Best Local Similarity
Matches 135; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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US-08-508-448C-25

Sequence 25, Application US/08508448C

Patent No. 5804410

GENERAL INFORMATION:

APPLICANT: Kazuyoshi YAWAOKA et al.

TITLE OF INVENTION: NUCLEIC ACID SEQUENCE ENCODING

TITLE OF INVENTION: TRYPSIN-LIKE BNZYME AND PROCESS FOR PRODUCING THE ENZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14;
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 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            246 RKHTDVFNWKVRAGSDKLGSFPSL--AVAKIIIIEFNPMYPKDNDIALMKLQFPLTFSGT 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   186 LVSLHCLACGKSLKTPRVVGGBEASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           128 NFTEALAETACROMGYSSKPTFRAVEIGPDODLDVVBITENSQELRMRN--SSGPCLSGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            132 ---KSRIESVLROMINNS----GNLEINP--STRITSLTDOAAANWLINECGAGPDLI--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73 YRTLSGRIESLITKTFKESNLRNQFIRAHVAKLRQDGSGVRADVVMKFQFTRNNNGASM-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   33 PILIALLSLASIIIVVVLIKVILDKYYFLCGOPLHFIPRK--QLCDGELDCPL-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 418;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24.3%; Score 568; DB 1; L 32.4%; Pred. No. 1.2e-49; ive 79; Mismatches 149;
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                                                                                                                                                                                                                                                                                                                                                                #700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette, 3.5 inch, 1.44 COMPUTER: IBM COMPATIBLE OPERATING SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEB: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordberfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/508,448C
FILING DATE: July 28, 1995
                                                                                774 CALPNRPGVYARVPRFTEWI 793
                                                             CGGPSTPGVYTKVSAYLNWI 427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: 202-371-8850
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INPERMATION:
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Warren M. Cheek, REGISTRATION NUMBER: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             138; Conservative
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MEDIUM TYPE: Diskett
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US-08-508-448C-25
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Matches 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLSGSLVSLHC--LACGKSLKT----PRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILD 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHWVLTAAHC-FRKHTDVFNWKVRAGSDKLGSF--PSLAVAKIIIIEFNPMY---PKDND 288
                                                                                                                                                                                                                  349 IDSTRCNADDAYQGEVTEKAMCAGIPEGGVDTCQGDSGGPLMYQ-SDQWHVVGIVSWGYG 407
                                                                                                                                                                                                                                                                                         IALMKLQFPLTFSGTVRPICLFFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQV 348
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                                                                          536 CLEDSLILLQCNYKSCGKKLVTQEVSPKIVGGSDSREGAWPWVVALYFDDQQVCGASLVS 595
                                       CLSGSLVSLHC--LACGKSLKT----PRVVGGEEASVDSWPWQVSIQYDKQHVCGGSILD 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLEDSLILLOCNYKSCGKKLVTQEVSPKIVGGSDSREGAMPWVALYFDDQQVCGASLVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      69 IPRKOLCDGELDCPLGEDEEHCVKSFPEGPAVAVRLSKDRSTLQVLDSATGNWFSACFDN
                                                                                                                                                                                        IALMKLQFPLTFSGTVRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: CLONING OF ENTEROKINASE AND METHOD OF USE NUMBER OF SEQUENCES: 33
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OFBRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
186 WITQISDDVCQLLGLGTGNSSVPTF-STGGGPYVNLNTAPNGSLILTPSQO-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24.4%; Score 571; DB 5; Length 798; 35.0%; Pred. No. 1.6e-49; ive 67; Mismatches 138; Indels
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APPLICATION NUMBER: PCT/US94/00616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application PC/TUS9400616
GENERAL INFORMATION:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                            CALPNRPGVYARVPRFTEWI 793
                                                                                                                                                                                                                                                                                                                                            CGGPSTPGVYTKVSAYLNWI 427
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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Matches 133;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           421
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                                                                                                                                                                                          Sequence 83, Application US/09370838;
Patent No. 6444425;
GENERAL INFORMATION:
APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadoh
APPLICANT: Secrist, Heather;
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF
TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.4750.
CURRENT FILING DATE: 1999-08-09
BARLIER FILING DATE: 1999-04-02
NUMBER OF SEQ ID NOS: 289
SOFTWARE: PSECSEQ for Windows Version 3.0
SEQ ID NO 83
LENGTH: 418
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32.4%; Pred. No. 1.2e-49;
ive 79; Mismatches 149; Indels
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APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Mohamath, Roadoh
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ORGANISM: Homo sapien
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APPLICANT: Reed, Steven G.
APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Lodes, Michael J.
APPLICANT: Monamath, Roadoh
APPLICANT: Secrist, Heather
TITLE OF INVENTION: LONG CANCER AND METHODS FOR THEIR USE
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TITLE OF INVENTION: LONG CANCER AND METHODS FOR THEIR USE
CURRENT APPLICATION NUMBER: US/09/370, 838
CURRENT FILING DATE: 1999-08-09
MUMBER OF SEQ ID NOS: 289
NUMBER OF SEQ ID NOS: 289
SOFTWARE: FastSEQ for Windows Version 3.0
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Patent No. 6444425
GENERAL INFORMATION:
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US-09-370-838-82
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Query Match 24.2%; Score 566; DB 4; Length 418;
Best Local Similarity 32.2%; Pred. No. 2e-49;
Matches 137; Conservative 80; Mismatches 149; Indels 60; Gaps 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    186 LVSLHCLACGKSLKTPRVVGGEBASVDSWPWQVSIQYDKQHVCGGSILDPHWVLTAAHCF 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            128 NFTEALASTACROMGYSSKPTFRAVEIGPDODLDVVEITENSQELRMRN--SSGPCLSGS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    246 RKHTDVFNWKVRAGSDKLGSFPSL--AVAKILILEFNPMYPKDNDIALMKLQFPLTFSGT 303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          304 VRPICLPFFDEELTPATPLWIIGWGFTKQNGGKMSDILLQASVQVIDSTRCNADDAYQGE 363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  364 VTEKMMCAGIPEGGVDTCQGDSGGPLMYQSDQ--WHVVGIVSWGYGCGGPSTPGVYTKVS 421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              73 YRTLSGRIESLITKTFKESNLRNQFIRAHVAKLRQDGSGVRADVVMKFQFTRNNNGASM- 131
                                                                                                                                                                                                                                                                                                                                                                                              ----GEDBEHCVKSFPEG-----PAVAVRLSKD----RSTLQVLDSATGNWFSACFD 127
                                                                                                                                                                                                                                                                                                                                                                        33 PIIIALLSLASIIIVVVLIKVILDKYYFLCGQPLHFIPRK--QLCDGELDCPL-----
TITLE OF INVENTION: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE TITLE OF INVENTION: LUNG CANCER AND METHODS FOR THEIR USE CURRENT APPLICATION NUMBER: US/09/370, 838
CURRENT RILING DATE: 1999-08-09
EARLIER APPLICATION NUMBER: US 09/285,323
NUMBER OF SEQ ID NOS: 289
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search completed: December 2, 2003, 02:31:09
Job time : 33.7125 secs
                                                                                                                                                                          SEQ ID NO 62
LENGTH: 418
TYPE: PRT
CRGANISM: Homo sapien
US-09-370-838-62
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